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(54) Title: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS USING PEPTIDE AND NU-CLEIC ACID COMPOSITIONS

(57) Abstract: This invention uses our knowledge of the mechanisms by which antigen is recognized by T cells to identify and prepare human papillomavirus (HPV) epitopes, and to develop epitope-based vaccines directed towards HPV. More specifically, this application communicates our discovery of pharmaceutical compositions and methods of use in the prevention and treatment of HPV infection.

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INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS

I. BACKGROUND OF THE INVENTION

Human papillomavirus (HPV) is a member of the papillomaviridae, a group of small DNA viruses that infect a variety of higher vertebrates. More than 80 types of HPVs have been identified. Of these, more than 30 can infect the genital tract. Some types, generally types 6 and 11, may cause genital wars, which are typically benign and rarely develop into cancer. Other strains of HPV, "cancer-associated", or "high-risk" types, can more frequently lead to the development of cancer. The primary mode of transmission of these strains of HPV is through sexual contact.

The main manifestations of the genital warts are cauliflower-like condylomata acuminata that usually involve moist surfaces; keratotic and smooth papular warts, usually on dry surfaces; and subclinical "flat" warts, which are found on any mucosal or cutaneous surface (Handsfield, H., Am. J. Med. 102(5A):16-20, 1997). These warts are typically benign but are a source of inter-individual spread of the virus (Ponten, J. & Guo, Z., Cancer Surv. 32:201-29, 1998). At least three HPV strains associated with genital warts have been identified: type 6a (see, e.g., Hofmann, K.J., et al., Virology 209(2):506-518, 1995), type 6b (see, e.g., Hofmann et al., supra) and type 11 (see, e.g., Dartmann, K. et al., Virology 151(1):124-130, 1986).

Cancer-associated HPVs have been linked with cancer in both men and women; they include, but are not limited to, HPV-16, HPV-18, HPV-31, HPV-45, HPV-33 and HPV-56. Other HPV strains, including types 6 and 11 as well as others, e.g., HPV-5 and HPV-8, are less frequently associated with cancer. The high risk types are typically associated with the development of cervical carcinoma and premalignant lesions of the cervix in women, but are also associated with similar malignant and premalignant lesions at other anatomic sites within the lower genital or anogenital tract. These lesions include neoplasia of the vagina, vulva, perineum, the penis, and the anus. HPV infection has also been associated with respiratory tract papillomas, and rarely, cancer, as well as abnormal growth or neoplasia in other epithelial tissues. See, e.g. VIROLOGY, 2²⁰⁰ ED, Fields et al., Eds. Raven Press, New York, 1990, Chapters 58 and 59, for a review of HPV association with cancer.

The HPV genome consists of three functional regions, the early region, the late region, and the "long control region". The early region gene products control viral replication, transcription and cellular transformation. They include the HPV E1 and E2 proteins, which play a role in HPV DNA replication, and the E6 and E7 oncoproteins, which are involved in the control of cellular proliferation. The late region include the genes that encode the structural proteins L1 and L2, which are the major and minor capsid proteins, respectively. The "long control region" contains such sequences as enhancer and promoter regulatory regions.

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HPV expresses different proteins at different stages of the infection, for example early, as well as late, proteins. Even in latent infections, however, early proteins are often expressed and are therefore useful targets for vaccine-based therapies. For example, high-grade dysplasia and cervical squamous cell carcinoma continue to express E6 and E7, which therefore can be targeted to treat disease at both early and late stages of infection.

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Treatment for HPV infection is often unsatisfactory because of persistence of virus after treatment and recurrence of clinically apparent disease is common. The treatment may require frequent visits to clinics and is not directed at elimination of the virus but at clearing warts. Because of persistence of virus after treatment, recurrence of clinically apparent disease is common.

Thus, a need exists for an efficacious vaccine to both prevent and treat HPV infection and to treat cancer that is associated with HPV infection. Effective HPV vaccines would be a significant advance in the control of sexually transmissable infections and could also protect against clinical disease, particularly cancers such as cervical cancer. (see, e.g., Rowen, P. & Lacey, C., Dermatologic Clinics 16(4):835-838, 1998).

Virus-specific, human leukocyte antigen (HLA) class I-restricted cytotoxic T lymphocytes (CTL) are known to play a major role in the prevention and clearance of virus infections in vivo (Oldstone et al., Nature 321:239, 1989, Jamieson et al., J. Virol. 61:3930, 1987; Yap et al., Nature 273:238, 1978; Lukacher et al., J. Exp. Med. 160:314, 1994; McMichael et al., N. Engl. J. Med. 309:13, 1983; Sethi et al., J. Gen. Virol. 64:443, 1983; Watari et al., J. Exp. Med. 165:459, 1987; Yasukawa et al., J. Immunol. 143:2051, 1989; Tigges et al., J. Virol. 66:1622, 1993; Reddenhase et al., J. Virol. 55:263, 1985; Quinnan et al., N. Engl. J. Med. 307:6, 1982). HLA class I molecules are expressed on the surface of almost all nucleated cells. Following intracellular processing of antigens, epitopes from the antigens are presented as a complex with the HLA class I molecules on the surface of such cells. CTL recognize the peptide-HLA class I complex, which then results in the destruction of the cell bearing the HLA-peptide complex directly by the CTL and/or via the activation of non-destructive mechanisms e.g., the production of interferon, that inhibit viral replication.

Virus-specific T helper lymphocytes are also known to be critical for maintaining effective immunity in chronic viral infections. Historically, HTL responses were viewed as primarily supporting the expansion of specific CTL and B cell populations; however, more recent data indicate that HTL may directly contribute to the control of virus replication. For example, a decline in CD4* T cells and a corresponding loss in HTL function characterize infection with HIV (Lane et al., New Engl. J. Med. 313:79, 1985). Furthermore, studies in HIV infected patients have also shown that there is an inverse relationship between virus-specific HTL responses and viral load, suggesting that HTL plays a role in viremia (see, e.g., Rosenberg et al., Science 278:1447, 1997).

The development of vaccines with prophylactic and therapeutic efficacy against HPV is ongoing. Early vaccine development was hampered by the inability to culture HPV. With the introduction of cloning techniques and protein expression, however, some attempts have been made to stimulate humoral and CTL response to HPV (See, e.g., Rowen, P. & Lacey, C., Dermatologic Clinics 16(4):335-838 (1998)) Studies to date, however, have been inconclusive.

Activation of T helper cells and cytotoxic lymphocytes (CTLs) in the development of vaccines has also been analyzed. Lehtinen, M., et al. for instance, has shown that some peptides from the E2 protein of HPV type 16 activate T helper cells and CTLs (Biochem. Biophys. Res. Commun. 209(2):541-6 (1995). Similarly, Tarpey et al, has shown that some peptides from HPV type 11 E7 protein can stimulate human HPV-specific CTLs in vitro (Immunology 81:222-227 (1994)) and Borysiewicz et al. have reported a recombinant vaccinia virus expressing HPV 16 and HPV 17 E6 and E7 that stimulated CTL responses in at least one patient (Lancet 347:1347-1357, 1996).

The epitope approach, as we have described, allows the incorporation of various antibody, CTL and HTL epitopes, from various proteins, in a single vaccine composition. Such a composition may simultaneously target multiple dominant and subdominant epitopes and thereby be used to achieve effective immunization in a diverse population.

The information provided in this section is intended to disclose the presently understood state of the art as of the filing date of the present application. Information is included in this section which was generated subsequent to the priority date of this application. Accordingly, information in this section is not intended, in any way, to delineate the priority date for the invention.

II. SUMMARY OF THE INVENTION

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This invention applies our knowledge of the mechanisms by which antigen is recognized by T cells, for example, to develop epitope-based vaccines directed towards HPV. More specifically, this application communicates our discovery of specific epitope pharmaceutical compositions and methods of use in the prevention and treatment of HPV infection.

Upon development of appropriate technology, the use of epitope-based vaccines has several advantages over current vaccines, particularly when compared to the use of whole antigens in vaccine compositions. There is evidence that the immune response to whole antigens is directed largely toward variable regions of the antigen, allowing for immune escape due to mutations. The epitopes for inclusion in an epitope-based vaccine may be selected from conserved regions of viral or tumor-associated antigens, which thereby reduces the likelihood of escape mutants. Furthermore, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of epitope-based vaccines.

An additional advantage of an epitope-based vaccine approach is the ability to combine selected epitopes (CTL and HTL), and further, to modify the composition of the epitopes, achieving, for example, enhanced immunogenicity. Accordingly, the immune response can be modulated, as appropriate, for the target disease. Similar engineering of the response is not possible with traditional approaches.

Another major benefit of epitope-based immune-stimulating vaccines is their safety. The possible pathological side effects caused by infectious agents or whole protein antigens, which might have their own intrinsic biological activity, is eliminated.

An epitope-based vaccine also provides the ability to direct and focus an immune response to multiple selected antigens from the same pathogen. Thus, patient-by-patient variability in the immune response to a particular pathogen may be alleviated by inclusion of epitopes from multiple antigens from the pathogen in a vaccine composition. In the case of HPV, epitopes derived from multiple strains may also be included. A "pathogen" may be an infectious agent or a tumor associated molecule.

One of the most formidable obstacles to the development of broadly efficacious epitopebased immunotherapeutics, however, has been the extreme polymorphism of HLA molecules. To date, effective non-genetically biased coverage of a population has been a task of considerable complexity; such coverage has required that epitopes be used that are specific for HLA molecules corresponding to each individual HLA allele. Impractically large numbers of epitopes would therefore have to be used in order to cover ethnically diverse populations. Thus, there has existed a need for peptide epitopes that are bound by multiple HLA antigen molecules for use in epitope-based vaccines. The greater the number of HLA antigen molecules bound, the greater the breadth of population coverage by the vaccine.

Furthermore, as described herein in greater detail, a need has existed to modulate peptide

binding properties, e.g., so that peptides that are able to bind to multiple HLA antigens do so with an

affinity that will stimulate an immune response. Identification of epitopes restricted by more than one HLA

allel at an affinity that correlates with immunogenicity is important to provide thorough population

coverage, and to allow the elicitation of responses of sufficient vigor to prevent or clear an infection in a

diverse segment of the population. Such a response can also target a broad array of epitopes. The

feelinology disclosed herein provides for such favored immune responses.

In a preferred embodiment, epitopes for inclusion in vaccine compositions of the invention are selected by a process whereby protein sequences of known antigens are evaluated for the presence of motif or supermotif-bearing epitopes. Peptides corresponding to a motif- or supermotif-bearing epitope are then synthesized and tested for the ability to bind to the HLA molecule that recognizes the selected motif. Those peptides that bind at an intermediate or high affinity i.e., an IC₃₀ (or a K₀ value) of 500 nM or less for HLA class I molecules or an IC₃₀ of 1000 nM or less for HLA class II molecules, are further evaluated for their ability to induce a CTL or HTL response. Immunogenic peptide epitopes are selected for inclusion in vaccine compositions.

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Supermotif-bearing peptides may additionally be tested for the ability to bind to multiple
alleles within the HLA supertype family. Moreover, peptide epitopes may be analogued to modify binding
affinity and/or the ability to bind to multiple alleles within an HLA supertype.

The invention also includes embodiments comprising methods for monitoring or evaluating an immune response to HPV in a patient having a known HLA-type. Such methods comprise incubating a T lymphocyte sample from the patient with a peptide composition comprising an HPV epitope that has an amino acid sequence described in Tables VII to Table XX which binds the product of at least one HLA allele present in the patient, and detecting for the presence of a T lymphocyte that binds to the peptide. A CTL peptide epitope may, for example, be used as a component of a tetrameric complex for this type of analysis.

An alternative modality for defining the peptide epitopes in accordance with the invention is to recite the physical properties, such as length; primary structure; or charge, which are correlated with binding to a particular allele-specific HLA molecule or group of allele-specific HLA molecules. A further modality for defining peptide epitopes is to recite the physical properties of an HLA binding pocket, or properties shared by several allele-specific HLA binding pockets (e.g. pocket configuration and charge distribution) and reciting that the peptide epitope fits and binds to the pocket or pockets.

As will be apparent from the discussion below, other methods and embodiments are also contemplated. Further, novel synthetic peptides produced by any of the methods described herein are also part of the invention.

5 III. DETAILED DESCRIPTION OF THE INVENTION

The peptides and corresponding nucleic acid compositions of the present invention are useful for stimulating an immune response to HPV by stimulating the production of CTL or HTL responses. The peptide epitopes, which are derived directly or indirectly from native HPV protein anino acid sequences, are able to bind to HLA molecules and stimulate an immune response to HPV. The complete sequence of the HPV proteins to be analyzed can be obtained from Genbank. Epitopes and analogs thereof can also be readily determined from sequence information that may subsequently be discovered for heretofore unknown variants of HPV, as will be clear from the disclosure provided below.

The epitopes of the invention have been identified in a number of ways, as will be discussed below. Also discussed in greater detail is that analog peptides have been derived and the binding 15 activity for HLA molecules modulated by modifying specific amino acid residues to create peptide analogs exhibiting altered immunogenicity. Further, the present invention provides compositions and combinations of compositions that enable epitope-based vaccines that are capable of interacting with HLA molecules encoded by various genetic alleles to provide broader population coverage than prior vaccines.

20 III.A. Definitions

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The invention can be better understood with reference to the following definitions, which are listed alphabetically:

A "computer" or "computer system" generally includes: a processor; at least one information storage/retrieval apparatus such as, for example, a hard drive, a disk drive or a tape drive; at least one input apparatus such as, for example, a keyboard, a mouse, a touch screen, or a microphone; and display structure. Additionally, the computer may include a communication channel in communication with a network. Such a computer may include more or less than what is listed above.

A "construct" as used herein generally denotes a composition that does not occur in nature. A construct can be produced by synthetic technologies, e.g., recombinant DNA preparation and expression or chemical synthetic techniques for nucleic or amino acids. A construct can also be produced by the addition or affiliation of one material with another such that the result is not found in nature in that form

"Cross-reactive binding" indicates that a peptide is bound by more than one HLA molecule; a synonym is degenerate binding.

A "cryptic epitope" elicits a response by immunization with an isolated peptide, but the response is not cross-reactive in vitro when intact whole protein which comprises the epitope is used as an antien.

A "dominant epitope" is an epitope that induces an immune response upon immunization with a whole native antigen (see, e.g., Sercarz, et al., Annu. Rev. Immunol. 11:729-766, 1993). Such a response is cross-reactive in vitro with an isolated peptide epitope.

With regard to a particular amino acid sequence, an "epitope" is a set of amino acid residues which is involved in recognition by a particular immunoglobulin, or in the context of T cells, those residues necessary for recognition by T cell receptor proteins and/or Major Histocompatibility Complex (MHC) receptors. In an immune system setting, in vivo or in vitro, an epitope is the collective features of a molecule, such as primary, secondary and tertiary peptide structure, and charge, that together form a site recognized by an immunoglobulin, T cell receptor or HLA molecule. Throughout this disclosure epitope and peptide are often used interchangeably. It is to be appreciated, however, that isolated or purified protein or peptide molecules larger than and comprising an epitope of the invention are still within the hounds of the invention.

"Human Leukocyte Antigen" or "HLA" is a human class I or class II Major
Histocompatibility Complex (MHC) protein (see, e.g., Stites, et al., IMMUNOLOGY, 8" ED., Lange
Publishing, Los Altos, CA (1994).

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An "HLA supertype or family", as used herein, describes sets of HLA molecules grouped on the basis of shared peptide-binding specificities. HLA class I molecules that share somewhat similar binding affinity for peptides bearing certain amino acid motifs are grouped into HLA supertypes. The terms HLA superfamily, HLA supertype family, HLA family, and HLA xx-like molecules (where xx denotes a particular HLA type), are synonyms.

Throughout this disclosure, results are expressed in terms of "IC₂₀'s." IC₂₀ is the concentration of peptide in a binding assay at which 50% inhibition of binding of a reference peptide is observed. Given the conditions in which the assays are run (i.e., limiting HLA proteins and labeled peptide concentrations), these values approximate K₀ values. Assays for determining binding are described in detail, e.g., in PCT publications WO 94/20127 and WO 94/03205. It should be noted that IC₂₀ values can change, often dramatically, if the assay conditions are varied, and depending on the particular reagents used (e.g., HLA preparation, etc.). For example, excessive concentrations of HLA molecules will increase the anaparent measured IC₂₀ of a given ligand.

Alternatively, binding is expressed relative to a reference peptide. Although as a particular assay becomes more, or less, sensitive, the IC₂₀'s of the peptides tested may change somewhat, the binding relative to the reference peptide will not significantly change. For example, in an assay run under conditions such that the IC₂₀ of the reference peptide increases 10-fold, the IC₂₀ values of the test peptides will also shift approximately 10-fold. Therefore, to avoid ambiguities, the assessment of whether a peptide is a good, intermediate, weak, or negative binder is generally based on its IC₅₀, relative to the IC₅₀ of a standard peptide.

Binding may also be determined using other assay systems including those using: live cells (e.g., Ceppellini et al., Nature 333-392, 1989; Christnick et al., Nature 352:67, 1991; Busch et al., Int. Immunol. 2:443, 19990; Hill et al., J. Immunol. 147:189, 1991; del Guercio et al., J. Immunol. 154:685, 1995), cell free systems using detergent lysates (e.g., Cerundol et al., J. Immunol. 2:12:069, 1991), immobilized purified MHC (e.g., Hill et al., J. Immunol. 152, 2890, 1994; Marshall et al., J. Immunol. 152:4946, 1994), ELISA systems (e.g., Reay et al., EMBO J. 11:2829, 1992), surface plasmon resonance (e.g., Khilko et al., J. Biol. Chem. 268:15425, 1993); high flux soluble phase assays (Hammer et al., J. Exp. Med. 180:2353, 1994), and measurement of class I MHC stabilization or assembly (e.g., Ljunggren et al., J. Exp.

Nature 346:476, 1990; Schumacher et al., Cell 62:563, 1990; Townsend et al., Cell 62:285, 1990; Parker et al., L. Immunol. 149:1896, 1992).

As used herein, "high affinity" with respect to HLA class I molecules is defined as binding with an IC₅₈ or K₀ value, of 50 nM or less, "intermediate affinity" is binding with an IC₅₈ or K₀ value of between about 500 and about 500 nM. "High affinity" with respect to binding to HLA class II molecules is defined as binding with an IC₅₈ or K₀ value of 100 aM or less; "intermediate affinity" is binding with an IC₅₈ or K₀ value of between about 100 and about 1000 nM.

The terms "identical" or percent "identity," in the context of two or more peptide sequences, refer to two or more sequences or subsequences that are the same or have a specified percentage of amino acid residues that are the same, when compared and aligned for maximum correspondence over a comparison window, as measured using a sequence comparison algorithm or by manual alignment and visual inspection.

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An "immunogenic peptide" or "peptide epitope" is a peptide that comprises an allelespecific motif or supermotif such that the peptide will bind an HLA molecule and induce a CTL and/or
15 HTL response. Thus, immunogenic peptides of the invention are capable of binding to an appropriate HLA
molecule and thereafter inducing a cytotoxic T cell response, or a helper T cell response, to the antigen
from which the immunogenic peptide is derived.

The phrases "isolated" or "biologically pure" refer to material which is substantially or essentially free from components which normally accompany the material as it is found in its native state. Thus, isolated peptides in accordance with the invention preferably do not contain materials normally associated with the peptides in their in situ environment.

"Link" or "join" refers to any method known in the art for functionally connecting peptides, including, without limitation, recombinant fusion, covalent bonding, disulfide bonding, ionic bonding, hydrogen bonding, and electrostatic bonding.

"Major Histocompatibility Complex" or "MHC" is a cluster of genes that plays a role in control of the cellular interactions responsible for physiologic immune responses. In humans, the MHC complex is also known as the HLA complex. For a detailed description of the MHC and HLA complexes, see, Paul, FINDAMENTAL IMMUNOLOGY, 3"6 ED., Raven Press, New York, 1993.

The term "motif" refers to the pattern of residues in a peptide of defined length, usually a

peptide of from about 8 to about 13 amino acids for a class I HLA motif and from about 6 to about 25

amino acids for a class II HLA motif, which is recognized by a particular HLA molecule. Peptide motifs

are typically different for each protein encoded by each human HLA allele and differ in the pattern of the

nrimary and secondary anchor residues.

A "negative binding residue" or "deleterious residue" is an amino acid which, if present at certain positions (typically not primary anchor positions) in a peptide epitope, results in decreased binding affinity of the peptide for the peptide's corresponding HLA molecule.

A "non-native" sequence or "construct" refers to a sequence that is not found in nature, i.e., is "non-naturally occurring". Such sequences include, e.g., peptides that are lipidated or otherwise modified, and polyepitopic compositions that contain epitopes that are not contiguous in a native protein sequence.

The term "peptide" is used interchangeably with "oligopeptide" in the present specification to designate a series of residues, typically L-amino acids, connected one to the other, typically by peptide bonds between the α -amino and carboxyl groups of adjacent amino acids. The preferred CTL-inducing peptides of the invention are 13 residues or less in length and usually consist of between about 8 and about 11 residues, preferably 9 or 10 residues. The preferred HTL-inducing oligopeptides are less than about 50 residues in length and usually consist of between about 6 and about 30 residues, more usually between about 12 and 25, and often between about 15 and 20 residues.

It is to be appreciated that protein or peptide molecules that comprise an epitope of the invention as well as additional amino acid(s) are within the bounds of the invention. In certain embodiments, there is a limitation on the length of a peptide of the invention which is not otherwise a construct as defined herein. An embodiment that is length-limited occurs when the protein/peptide comprising an epitope of the invention comprises a region (i.e., a contiguous series of amino acids) having 100% identity with a native sequence. In order to avoid a recited definition of epitope from reading, e.g., on whole natural molecules, the length of any region that has 100% identity with a native peptide sequence is limited. Thus, for a peptide comprising an epitope of the invention and a region with 100% identity with a native peptide sequence (and which is not otherwise a construct), the region with 100% identity to a native sequence generally has a length of: less than or equal to 600 amino acids, often less than or equal to 500 amino acids, often less than or equal to 400 amino acids, often less than or equal to 250 amino acids. often less than or equal to 100 amino acids, often less than or equal to 85 amino acids, often less than or equal to 75 amino acids, often less than or equal to 65 amino acids, and often less than or equal to 50 amino acids. In certain embodiments, an "epitope" of the invention which is not a construct is comprised by a peptide having a region with less than 51 amino acids that has 100% identity to a native peptide sequence, in any increment of (50, 49, 48, 47, 46, 45, 44, 43, 42, 41, 40, 39, 38, 37, 36, 35, 34, 33, 32, 31, 30, 29, 28, 27, 26, 25, 24, 23, 22, 21, 20, 19, 18, 17, 16, 15, 14, 13, 12, 11, 10, 9, 8, 7, 6, 5) down to 5 amino acids.

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Certain peptide or protein sequences longer than 600 amino acids are within the scope of the invention. Such longer sequences are within the scope of the invention so long as they do not comprise any contiguous sequence of more than 600 amino acids that have 100% identity with a native peptide sequence, or if longer than 600 amino acids, they are a construct. For any peptide that has five contiguous residues or less that correspond to a native sequence, there is no limitation on the maximal length of that peptide in order to fall within the scope of the invention. It is presently preferred that a CTL epitope of the invention be less than 600 residues long in any increment down to eight amino acid residues.

"Pharmaceutically acceptable" refers to a non-toxic, inert, and/or physiologically compatible composition.

A "pharmaceutical excipient" comprises a material such as an adjuvant, a carrier, pHadjusting and buffering agents, tonicity adjusting agents, wetting agents, preservative, and the like.

A "primary anchor residue" is an amino acid at a specific position along a peptide sequence which is understood to provide a contact point between the immunogenic peptide and the HLA molecule. One to three, usually two, primary anchor residues within a peptide of defined length generally defines a "motif" for an immunogenic peptide. These residues are understood to fit in close contact with peptide binding grooves of an HLA molecule, with their side chains buried in specific pockets of the

binding grooves themselves. In one embodiment, for example, the primary anchor residues are located at position 2 (from the amino terminal position) and at the carboxyl terminal position of a 9-residue peptide epitope in accordance with the invention. The primary anchor positions for each motif and supermotif are set forth in Table 1. For example, analog peptides can be created by altering the presence or absence of particular residues in these primary anchor positions. Such analogs are used to modulate the binding affinity of a peptide comprising a particular motif or supermotif.

"Promiscuous recognition" is where a distinct peptide is recognized by the same T cell clone in the context of various HLA molecules. Promiscuous recognition or binding is synonymous with cross-reactive binding.

A "protective immune response" or "therapeutic immune response" refers to a CTL and/or an HTL response to an antigen derived from an infectious agent or a tumor antigen, which prevents or at least partially arrests disease symptoms or progression. The immune response may also include an antibody response which has been facilitated by the stimulation of helper T cells.

The term "residue" refers to an amino acid or amino acid mimetic incorporated into an oligopeptide by an amide bond or amide bond mimetic.

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A "secondary anchor residue" is an amino acid at a position other than a primary anchor position in a peptide which may influence peptide binding. A secondary anchor residue occurs at a significantly higher frequency amongst bound peptides than would be expected by random distribution of amino acids at one position. The secondary anchor residues are said to occur at "secondary anchor positions." A secondary anchor residue can be identified as a residue which is present at a higher frequency among high or intermediate affinity binding peptides, or a residue otherwise associated with high or intermediate affinity binding. For example, analog peptides can be created by altering the presence or absence of particular residues in these secondary anchor positions. Such analogs are used to finely modulate the binding affinity of a peptide comprising a particular motif of supermotif.

A "subdominant epitope" is an epitope which evokes little or no response upon immunization with whole antigens which comprise the epitope, but for which a response can be obtained by immunization with an isolated peptide, and this response (unlike the case of cryptic epitopes) is detected when whole protein is used to recall the response in vitro or in vivo.

A "supermotif" is a peptide binding specificity shared by HLA molecules encoded by two or more HLA alleles. Preferably, a supermotif-bearing peptide is recognized with high or intermediate affinity (as defined herein) by two or more HLA antigens.

"Synthetic peptide" refers to a peptide that is man-made using such methods as chemical synthesis or recombinant DNA technology.

As used herein, a "vaccine" is a composition that contains one or more peptides of the invention. There are numerous embodiments of vaccines in accordance with the invention, such as by a cocktail of one or more peptides; one or more epitopes of the invention comprised by a polyepitopic peptide; or nucleic acids that encode such peptides or polypeptides, e.g., a minigene that encodes a polyepitopic peptide. The "one or more peptides" can include any whole unit integer from 1-150, e.g., at least 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 105, 110, 115, 120, 125,

130, 135, 140, 145, or 150 or more peptides of the invention. The peptides or polypeptides can optionally be modified, such as by lipidation, addition of targeting or other sequences. HLA class I-binding peptides of the invention can be admixed with, or linked to, HLA class II-binding peptides, to facilitate activation of both cytotoxic T lymphocytes and helper T lymphocytes. Vaccines can also comprise peptide-pulsed antigen presenting cells, e.g., dendritic cells.

The nomenclature used to describe peptide compounds follows the conventional practice wherein the amino group is presented to the left (the N-terminus) and the carboxyl group to the right (the Cterminus) of each amino acid residue. When amino acid residue positions are referred to in a peptide epitope they are numbered in an amino to carboxyl direction with position one being the position closest to the amino terminal end of the epitope, or the peptide or protein of which it may be a part. In the formulae representing selected specific embodiments of the present invention, the amino- and carboxyl-terminal groups, although not specifically shown, are in the form they would assume at physiologic pH values, unless otherwise specified. In the amino acid structure formulae, each residue is generally represented by standard three letter or single letter designations. The L-form of an amino acid residue is represented by a capital single letter or a capital first letter of a three-letter symbol, and the D-form for those amino acids having D-forms is represented by a lower case single letter or a lower case three letter symbol. Glycine has no asymmetric carbon atom and is simply referred to as "Gly" or G. The amino acid sequences of peptides set forth herein are generally designated using the standard single letter symbol. (A, Alanine; C, Cysteine; D. Aspartic Acid; E, Glutamic Acid; F, Phenylalanine; G, Glycine; H, Histidine; I, Isoleucine; K, Lysine; L, Leucine; M, Methionine; N, Asparagine; P, Proline; Q, Glutamine; R, Arginine; S, Serine; T, Threonine; V, Valine; W, Tryptophan; and Y, Tyrosine.) In addition to these symbols, "B" in the single letter

abbreviations used herein designates α-amino butyric acid.

III.B. Stimulation of CTL and HTL responses

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The mechanism by which T cells recognize antigens has been delineated during the past ten years. Based on our understanding of the immune system we have developed efficacious pentide epitope vaccine compositions that can induce a therapeutic or prophylactic immune response to HPV in a broad population. For an understanding of the value and efficacy of the claimed compositions, a brief review of immunology-related technology is provided.

A complex of an HLA molecule and a peptidic antigen acts as the ligand recognized by HLA-restricted T cells (Buus, S. et al., Cell 47:1071, 1986; Babbitt, B. P. et al., Nature 317:359, 1985; Townsend, A. and Bodmer, H., Annu. Rev. Immunol. 7:601, 1989; Germain, R. N., Annu. Rev. Immunol. 11:403, 1993). Through the study of single amino acid substituted antigen analogs and the sequencing of endogenously bound, naturally processed peptides, critical residues that correspond to motifs required for specific binding to HLA antigen molecules have been identified and are described herein and are set forth in Tables I, II, and III (see also, e.g., Southwood, et al., J. Immunol. 160:3363, 1998; Rammensee, et al., Immunogenetics 41:178, 1995; Rammensee et al., SYFPEITHI, access via web at: http://134.2.96.221/scripts.hlaserver.dll/home.htm; Sette, A. and Sidney, J. Curr. Opin. Immunol. 10:478, 1998; Engelhard, V. H., Curr. Opin. Immunol. 6:13, 1994; Sette, A. and Grey, H. M., Curr. Opin. Immunol. 4:79, 1992; Sinigaglia, F. and Hammer, J. Curr. Biol. 6:52, 1994; Ruppert et al., Cell 74:929-937, 1993;

Kondo et al., J. Immunol. 155:4307-4312, 1995, Sidney et al., J. Immunol. 157:3480-3490, 1996; Sidney et al., Human Immunol. 45:79-93, 1996; Sette, A. and Sidney, J. Immunogenetics 1999 Nov. 50(3-4):201-12.
Review).

Furthermore, x-ray crystallographic analysis of HLA-peptide complexes has revealed

5 pockets within the peptide binding cleft of HLA molecules which accommodate, in an allele-specific mode, residues borne by peptide ligands; these residues in turn determine the HLA binding capacity of the peptides in which they are present. (See, e.g., Madden, D.R. Annu. Rev. Immunol. 13:587, 1995; Smith, et al., Immunity 4:203, 1996; Fremont et al., Immunity 8:305, 1998; Stern et al., Structure 2:245, 1994; Jones, E.Y. Curr. Opin. Immunol. 9:75, 1997; Brown, J. H. et al., Nature 364:33, 1993; Guo, H. C. et al., Proc.

10 Natl. Acad. Sci. USA 90:8053, 1993; Guo, H. C. et al., Nature 360:364, 1992; Silver, M. L. et al., Nature 360:367, 1992; Matsumura, M. et al., Science 257:927, 1992; Madden et al., Cell 70:1035, 1992; Fremont, D. H. et al., Science 257:919, 1992; Saper, M. A., Bjorkman, P. J. and Wiley, D. C., J. Mol. Biol. 219:277, 1991.)

Accordingly, the definition of class I and class II allele-specific HLA binding motifs, or

15 class I or class II supermotifs allows identification of regions within a protein that have the potential of binding particular HLA antigen(s).

The present inventors have found that the correlation of binding affinity with immunogenicity, which is disclosed herein, is an important factor to be considered when evaluating candidate peptides. Thus, by a combination of motif searches and HLA-peptide binding assays, candidates for epitope-based vaccines have been identified. After determining their binding affinity, additional confirmatory work can be performed to select, amongst these vaccine candidates, epitopes with preferred characteristics in terms of population coverage, antigenicity, and immunogenicity.

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Various strategies can be utilized to evaluate immunogenicity, including:

1) Evaluation of primary T cell cultures from normal individuals (see, e.g., Wentworth, P.

- 25 A. et al., Mol. Immunol. 32:603, 1995; Celis, E. et al., Proc. Natl. Acad. Sci. USA 91:2105, 1994; Tsai, V. et al., J. Immunol. 158:1796, 1997; Kawashima, I. et al., Human Immunol. 59:1, 1998); This procedure involves the stimulation of peripheral blood lymphocytes (PBL) from normal subjects with a test peptide in the presence of antigen presenting cells in vitro over a period of several weeks. T cells specific for the peptide become activated during this time and are detected using, e.g., a lymphokine- or 51Cr-release assay involving peptide sensitized target cells.
 - 2) Immunization of HLA transgenic mice (see, e.g., Wentworth, P. A. et al., J. Immunol. 26:97, 1996; Wentworth, P. A. et al., Int. Immunol. 8:651, 1996; Alexander, J. et al., J. Immunol. 159:4753, 1997); In this method, peptides in incomplete Freund's adjuvant are administered subcutaneously to HLA transgenic mice. Several weeks following immunization, splenocytes are removed and cultured in vitro in the presence of test peptide for approximately one week. Peptide-specific T cells are detected using, e.g., a 51Cr-release assay involving peptide sensitized target cells and target cells expressing endogenously generated antigen.
 - 3) Demonstration of recall T cell responses from immune individuals who have effectively been vaccinated, recovered from infection, and/or from chronically infected patients (see, e.g., Rehermann, B. et al., J. Exp. Med. 181:1047, 1995; Doolan, D. L. et al., Immunity 7:97, 1997; Bertoni, R.

et al., J. Clin. Invest. 100:503, 1997; Threlkeld, S. C. et al., J. Immunol. 159:1648, 1997; Diepolder, H. M. et al., J. Virol. 71:6011, 1997). In applying this strategy, recall responses are detected by culturing PBL from subjects that have been naturally exposed to the antigen, for instance through infection, and thus have generated an immune response "naturally", or from patients who were vaccinated against the infection. PBL from subjects are cultured in vitro for 1-2 weeks in the presence of test peptide plus antigen presenting cells (APC) to allow activation of "memory" T cells, as compared to "naive" T cells. At the end of the culture period, T cell activity is detected using assays for T cell activity including 51Cr release involving peptide-sensitized targets, T cell proliferation, or lymphokine release.

The following describes the peptide epitopes and corresponding nucleic acids of the invention.

III.C. Binding Affinity of Peptide Epitopes for HLA Molecules

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As indicated herein, the large degree of HLA polymorphism is an important factor to be taken into account with the epitope-based approach to vaccine development. To address this factor, epitope selection encompassing identification of peptides capable of binding at high or intermediate affinity to multiple HLA molecules is preferably utilized, most preferably these epitopes bind at high or intermediate affinity to two or more allele-specific HLA molecules.

CTL-inducing peptides of interest for vaccine compositions preferably include those that have an IC₅₉ or binding affinity value for class I HLA molecules of 500 nM or better (i.e., the value is ≤ 500 nM). HTL-inducing peptides preferably include those that have an IC₅₉ or binding affinity value for class II HLA molecules of 1000 nM or better, (i.e., the value is ≤ 1,000 nM). For example, peptide binding is assessed by testing the capacity of a candidate peptide to bind to a purified HLA molecule in vitro. Peptides exhibiting high or intermediate affinity are then considered for further analysis. Selected peptides are tested on other members of the supertype family. In preferred embodiments, peptides that exhibit cross-reactive binding are then used in cellular screening analyses or vaccines.

As disclosed herein, higher HLA binding affinity is correlated with greater immunogenicity. Greater immunogenicity can be manifested in several different ways. Immunogenicity corresponds to whether an immune response is elicited at all, and to the vigor of any particular response, as well as to the extent of a population in which a response is elicited. For example, a peptide might elicit an immune response in a diverse array of the population, yet in no instance produce a vigorous response. In accordance with these principles, close to 90% of high binding peptides have been found to be immunogenic, as contrasted with about 50% of the peptides which bind with intermediate affinity. Moreover, higher binding affinity peptides lead to more vigorous immunogenic responses. As a result, less peptide is required to elicit a similar biological effect if a high affinity binding peptide is used. Thus, in preferred embodiments of the invention, high affinity binding pitopes are particularly useful.

The relationship between binding affinity for HLA class I molecules and immunogenicity of discrete peptide epitopes on bound antigens has been determined for the first time in the art by the present inventors. The correlation between binding affinity and immunogenicity was analyzed in two different experimental approaches (see, e.g., Sette, et al., J. Immunol. 153:5586-5592, 1994). In the first approach, the immunogenicity of potential epitopes ranging in HLA binding affinity over a 10,000-fold

range was analyzed in HLA-A*0201 transgenic mice. In the second approach, the antigenicity of approximately 100 different hepatitis B virus (HBV)-derived potential epitopes, all carrying A*0201 binding motifs, was assessed by using PBL from acute hepatitis patients. Pursuant to these approaches, it was determined that an affinity threshold value of approximately 500 aM (preferably 50 nM or less) determines the capacity of a peptide epitope to elicit a CTL response. These data are true for class I binding affinity measurements for naturally processed peptides and for synthesized T cell epitopes. These data also indicate the important role of determinant selection in the shaping of T cell responses (see, e.g., Schaefter et al. Proc. Natl. Acad. Sci. USA 86:4649-4653, 1989).

An affinity threshold associated with immunogenicity in the context of HLA class II DR molecules has also been delineated (see, e.g., Southwood et al. J. Immunology 160:3363-3373,1998, and co-pending U.S.S.N. 09/009,953 filed 1/21/98). In order to define a biologically significant threshold of DR binding affinity, a database of the binding affinities of 32 DR-restricted epitopes for their restricting element (i.e., the HLA molecule that binds the motif) was compiled. In approximately half of the cases (15 of 32 epitopes), DR restriction was associated with high binding affinities, i.e. binding affinity values of 100 nM or less. In the other half of the cases (16 of 32), DR restriction was associated with intermediate affinity (binding affinity values in the 100-1000 nM range). In only one of 32 cases was DR restriction associated with an IC₂₉ of 1000 nM or greater. Thus, 1000 nM can be defined as an affinity threshold associated with immunogenicity in the context of DR molecules.

In the case of tumor-associated antigens (TAAs), many CTL peptide epitopes that have

been shown to induce CTL that lyse peptide-pulsed target cells and tumor cell targets endogenously
expressing the epitope exhibit binding affinity or 1C₉ values of 200 nM or less. In a study that evaluated
the association of binding affinity and immunogenicity of a small set of such TAA epitopes, 100% (10/10)
of the high binders, i.e., peptide epitopes binding at an affinity of 50 nM or less, were immunogenic and
80% (8/10) of them elicited CTLs that specifically recognized tumor cells. In the 51 to 200 nM range, very
similar figures were obtained. With respect to analog peptides, CTL inductions positive for wildtype
peptide and tumor cells were noted for 86% (6/7) and 71% (5/7) of the peptides, respectively. In the 201500 nM range, most peptides (4/5 wildtype) were positive for induction of CTL recognizing wildtype
peptide, but tumor recognition was not detected.

The binding affinity of peptides for HLA molecules can be determined as described in 30 Example 1, below.

III.D. Peptide Epitope Binding Motifs and Supermotifs

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Through the study of single amino acid substituted antigen analogs and the sequencing of endogenously bound, naturally processed peptides, critical residues required for allele-specific binding to HLA molecules have been identified. The presence of these residues correlates with binding affinity for HLA molecules. The identification of motifs and/or supermotifs that correlate with high and intermediate affinity binding is an important issue with respect to the identification of immunogenic peptide epitopes for the inclusion in a vaccine. Kast et al. (J. Immunol. 152:3904-3912, 1994) have shown that motif-bearing peptides account for 90% of the epitopes that bind to allele-specific HLA class I molecules. In this study all possible peptides of 9 amino acids in length and overlapping by eight amino acids (240 peptides), which

cover the entire sequence of the E6 and E7 proteins of human papillomavirus type 16, were evaluated for binding to five allele-specific HLA molecules that are expressed at high frequency among different ethnic groups. This unbiased set of peptides allowed an evaluation of the predictive value of HLA class I motifs. From the set of 240 peptides, 22 peptides were identified that bound to an allele-specific HLA molecule with high or intermediate affinity. Of these 22 peptides, 20 (i.e. 91%) were motif-bearing. Thus, this study demonstrates the value of motifs for the identification of peptide epitopes for inclusion in a vaccine: application of motif-based identification techniques will identify about 90% of the potential epitopes in a target antigen protein sequence.

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Such peptide epitopes are identified in the Tables described below.

Peptides of the present invention may also comprise epitopes that bind to MHC class II DR molecules. A greater degree of heterogeneity in both size and binding frame position of the motif, relative to the N and C termini of the peptide, exists for class II peptide ligands. This increased heterogeneity of HLA class II peptide ligands is due to the structure of the binding groove of the HLA class II molecule which, unlike its class I counterpart, is open at both ends. Crystallographic analysis of HLA class II DRB*0101-peptide complexes showed that the major energy of binding is contributed by peptide residues complexed with complementary pockets on the DRB*0101 molecules. An important anchor residue engages the deepest hydrophobic pocket (see, e.g., Madden, D.R. Ann. Rev. Immunol. 13:587, 1995) and is referred to as position 1 (P1). P1 may represent the N-terminal residue of a class II binding peptide epitope, but more typically is flanked towards the N-terminaus by one or more residues. Other studies have also pointed to an important role for the peptide residue in the 6th position towards the C-terminus, relative to P1, for binding to various DR molecules.

In the past few years evidence has accumulated to demonstrate that a large fraction of HLA class I and class II molecules can be classified into a relatively few supertypes, each characterized by largely overlapping peptide binding repertoires, and consensus structures of the main peptide binding 25 pockets. Thus, peptides of the present invention are identified by any one of several HLA-specific amino acid motifs (see, e.g., Tables I-III), or if the presence of the motif corresponds to the ability to bind several allele-specific HLA antigens, a supermotif. The HLA molecules that bind to peptides that possess a particular amino acid supermotif are collectively referred to as an HLA "supertype."

The peptide motifs and supermotifs described below, and summarized in Tables I-III, provide guidance for the identification and use of peptide epitopes in accordance with the invention.

Examples of peptide epitopes bearing a respective supermotif or motif are included in Tables as designated in the description of each motif or supermotif below. The Tables include a binding affinity ratio listing for some of the peptide epitopes. The ratio may be converted to \mathbb{C}_{50} by using the following formula: \mathbb{C}_{50} of the standard peptide/ratio = \mathbb{C}_{50} of the test peptide (i.e., the peptide epitope). The \mathbb{C}_{50} values of standard peptides used to determine binding affinities for Class I peptides are shown in Table IV. The \mathbb{C}_{50} values of standard peptides used to determine binding affinities for Class II peptides are shown in Table V. For example, where an HLA-A.2.1 motif-bearing peptide shows a relative binding ratio of 0.01 for HLA-A-90201, the \mathbb{C}_{50} value is 500 nM, and where an HLA-A.2.1 motif-bearing peptide shows a relative binding ratio of 0.1 for HLA-A-90201, the \mathbb{C}_{50} value is 500 nM.

The peptides used as standards for the binding assays described herein are examples of standards; alternative standard peptides can also be used when performing binding studies.

To obtain the peptide epitope sequences listed in Tables VII-XX, protein sequence data for HPV types 6a, 6b, 11a, 16, 18, 31, 33, 45, and 56 were evaluated for the presence of the designated supermotif or motif. Seven HPV structural and regulatory proteins, E1, E2, E5, E6, E7, L1 and L2 were included in the analysis. E4 was also included in the evaluation of some of the strains. Peptide epitopes can additionally be evaluated on the basis of their conservancy (i.e., the amount of variance) among the available protein sequences for each HPV antigen.

In the Tables, motif- and/or supermotif-bearing amino acids sequences identified in the indicated HPV strains are designated by position number and length of the epitope with reference to the HPV sequences and numbering provided below. For each sequence, the four columns provide the following information: column I indicates the HPV strain; column 2 indicates the HPV protein in which the motif-bearing sequence is found, e.g., El, E2, E4, E5, E6, E7, L1, or L2; column 3 indicates the length of the epitope, or in the case of HLA Class II epitopes, the length of the core sequence; and column 4 designates the amino acid position in the HPV protein sequence that corresponds to the first amino acid residue of the epitope. For those sections of the Tables that include only three columns, corresponding to columns 2, 3, and 4, the HPV strain is indicated in the heading at the top of the page. For example, the first peptide epitope listed in Table VII, i.e., the HLA-A1 supermotif, for HPV 16, protein E1 is a sequence of 10 residues in length starting at position 206. Accordingly, the amino acid sequence of the epitope is

For HPV strain 11, the number and position listed for protein E5 refers to either the HPV11 E5a or HPV11 E5b sequence set out below. Because the epitope must include the designated motif or supermotif, e.g., HLA-A2, it can readily be determined whether the sequence refers to HPV11 E5a or E5b by checking the amino acid sequences of both E5a and E5b and selecting the sequence that conforms to the motif listed in Table I.

HPV STRAINS AND AMINO ACID SEQUENCES OF HPV PROTEINS

HPV6A E1

1 MADDSGTENEGSGCTGWFWVEAIVOHPTGTQISDDEDEVEDSGYDMVDFIDDSNITHNS 60
30 LEAQALFNRQEADTHYATVQDLKRKYLGSPYVSPINTIAEAVESEISPRLDAIKLTRQPK 120
KYKRRLFCTRELITDSGYGYSSEVEAGTGTQVEKHGVPENGGGCQEKDTGRDIEGEBHTEAE 120
APTINSVREHAGTAGILELLKCKDLRAALLGKFKECFGLSFIDLIRPFKSDKTTCADWVVA 240
GFGIHHSISEAFQKLIEPLSLYAHIQWLTNAWGMVLLVLVRFKVNISKSTVARTLATLLIN 300
1EDNQMLIEPPRIQSGVAALWFRTGISNASTVIGEAPEWITRQTVIEHGLADSQFKLTE 360
MYQMAYDNDICEESEIAFEYAQRGDFDSNARAFLNSNMQAKYVKDCATMCRHYKHAEMRK 420
MSIKQWIKHRGSKIBGTGNWKBIVOFLHHONLEFIPPLSKFKLMLHGTPKNGCIAIVGPP 480
DTGKSYFCMSLIFFLGGTVISHVNSSSHFMLQPLVDAKVALLDDATQPCWIYMDTYMRLL 540
LDGNPMSIDRKHKALTLIKCPPLLVTSNIDITKEEKYKYLHTRUTTFTPNPPFFDRNGN 600
AVYELSNAMWKCFFERLSSSLDIOGSEDESDGSNQAFRCVFGTVVTTL 649

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	HPV6A_E2	
	1 MEAIAKRLDACQEQLLELYEENSTDLNKHVLHWKCMRHESVLLYKAKQMGLSHIGMQVVP	60
	PLKVSEAKGHNAIEMQMHLESLLKTEYSMEPWTLQETSYEMWQTPPKRCFKKRGKTVEVK	120
	FDGCANNTMDYVVWTDVYVQDTDSWVKVHSMVDAKGIYYTCGQFKTYYVNFVKEAEKYGS	180
5	TKQWEVCYGSTVICSPASVSSTTQEVSIPESTTYTPAQTSTPVSSSTQEDAVQTPPRKRA	240
	RGVQQSPCNALCVAHIGPVDSGNHNLITNNHDQHQRRNNSNSSATPIVQFQGESNCLKCF	300
	RYRLNDKHRHLFDLISSTWHWASPKAPHKHAIVTVTYHSEEQRQQFLNVVKIPPTIRHKL	360
	GFMSLHLL 368	
10	HDV6A E4	
	1 MAAQLYVLLHLYLALHKKYPFLNLLHTPPHRPPPLCPQAPRKTQCKRRLENEHEESNSHL	60
	ATPCVWPTLDPWTVETTTSSLTITTSTKEGTTVTVQLRL 99	
	HPV6A E5	
15	1 MEVVPVQIAAGTTSTLILPVIIAFVVCFVSIILIVWISDFIVYTSVLVLTLLLYLLLWLL	60
	LTTPLQFFLLTLLVCYCPALYIHHYIVNTQQ 91	
	HPV6A E6	
	1 MESANASTSATTIDQLCKTFNLSMHTLQINCVFCKNALTTAEIYSYAYKQLKVLFRGGYP	60
20	YAACACCLEFHGKINQYRHFDYAGYATTVEEETKQDILDVLIRCYLCHKFLCEVEKVKHI	120
	LTKARFIKLNCTWKGRCLHCWTTCMEDMLP 150	
	HPV6A E7	
	1 MHGRHVTLKDIVLDLQPPDPVGLHCYEQLVDSSEDEVDEVDGQDSQPLKQHFQIVTCCCG	60
25	CDSNVRLVVQCTETDIREVQQLLLGTLDIVCPICAPKT 98	
	HPV6A L1	
	1 MWRPSDSTVYVPPPNPVSKVVATDAYVTRTNIFYHASSSRLLAVGHPYFSIKRANKTVVP	60
	KVSGYQYRVFKVVLPDPNKFALPDSSLFDPTTQRLVWACTGLEVGRGQPLGVGVSGHPFL	120
30	NKYDDVENSGSGGNPGQDNRVNVGMDYKQTQLCMVGCAPPLGEHWGKGKQCTNTPVQAGD	180
	${\tt CPPLELITSVIQDGDMVDTGFGAMNFADLQTNKSDVPIDICGTTCKYPDYLQMAADPYGD}$	240
	RLFFFLRKEQMFARHFFNRAGEVGEPVPDTLIIKGSGNRTSVGSSIYVNTPSGSLVSSEA	300
	QLFNKPYWLQKAQGHNNGICWGNQLFVTVVDTTRSTNMTLCASVTTSSTYTNSDYKEYMR	360
	HVEEYDLQFIFQLCSITLSAEVMAYIHTMNPSVLEDWNFGLSPPPNGTLEDTYRYVQSQA	420
35	ITCQKPTPEKEKPDPYKNLSFWEVNLKEKFSSELDQYPLGRKFLLQSGYRGRSSIRTGVK RPAVSKASAAPKRKRAKTKR 500	480
	UUC MIIAKAANAYAACAACVAYA	
	HPV6A L2 1 MAHSRARRRKRASATOLYQTCKLTGTCPPDVIPKVEHNTIADQILKWGSLGVFFGGLGIG	60
40	1 MAHSRARRRKRASATQLYQTCKLTGTCPPDVTFAVBARTTADQTBARGSBSVFFGGGGT TGSGTGGRTGYVPLGTSAKPSITSGPMARPPVVVEPVAPSDPSIVSLIEESAIINAGAPE	120
40	TGSGTGGRTGYVPLGTSAKPSITSGPMARPPVVVEPVAPSDFSIVSDFEBSATIANSH E	180

	SAPTITSHPIEEIPLDTFVISSSDSGPTSSTPVPGTAPRPRVGLYSRALHQVQVTDPAFL	240
	STPQRLITYDNPVYEGEDVSVQFSHDSIHNAPDEAFMDIIRLHRPAIASRRGLVRYSRIG	300
	QRGSMHTRSGKHIGARIHYFYDISPIAQAAEEIEMHPLVAAQDDTFDIYAESFEPDINPT	360
	QHPVTNISDTYLTSTPNTVTQPWGNTTVPLSSIPNDLFLQSGPDITFPTAPMGTPFSPVT	420
5	ALPTGPVFITGSGFYLHPAWYFARKRRKRIPLFFSDVAA 459	
	HPV6B E1	
	1 MADDSGTENEGSGCTGWFMVEAIVQHPTGTQISDDEDEEVEDSGYDMVDFIDDSNITHNS	60
	LEAQALFNRQEADTHYATVQDLKRKYLGSPYVSPINTIAEAVESEISPRLDAIKLTRQPK	120
10	KVKRRLFQTRELTDSGYGYSEVEAGTGTQVEKHGVPENGGDGQEKDTGRDIEGEEHTEAE	180
	APTNSVREHAGTAGILELLKCKDLRAALLGKFKECFGLSFIDLIRPFKSDKTTCLDWVVA	240
	GFGIHHSISEAFQKLIEPLSLYAHIQWLTNAWGMVLLVLLRFKVNKSRSTVARTLATLLN	300
	IPENQMLIEPPKIQSGVAALYWFRTGISNASTVIGEAPEWITRQTVIEHGLADSQPKLTE	360
	MVQWAYDNDICEESEIAFEYAQRGDFDSNARAFLNSNMQAKYVKDCATMCRHYKHAEMRK	420
15	MSIKQWIKHRGSKIEGTGNWKPIVQFLRHQNIEFIPFLTKFKLWLHGTPKKNCIAIVGPP	480
	DTGKSYFCMSLISFLGGTVISHVNSSSHFWLQPLVDAKVALLDDATQPCWIYMDTYMRNL	540
	LDGNPMSIDRKHKALTLIKCPPLLVTSNIDITKEDKYKYLHTRVTTFTFPNPFPFDRNGN	600
	AVYELSNTNWKCFFERLSSSLDIQDSEDEEDGSNSQAFRCVPGTVVRTL 649	
20	HPV6B E2	
	1 MEAIAKRLDACQEQLLELYEENSTDLHKHVLHWKCMRHESVLLYKAKQMGLSHIGMQVVP	60
	PLKVSEAKGHNAIEMQMHLESLLRTEYSMEPWTLQETSYEMWQTPPKRCFKKRGKTVEVK	120
	FDGCANNTMDYVVWTDVYVQDNDTWVKVHSMVDAKGIYYTCGQFKTYYVNFVKEAEKYGS	180
	TKHWEVCYGSTVICSPASVSSTTQEVSIPESTTYTPAQTSTLVSSSTKEDAVQTPPRKRA	240
25	RGVQQSPCNALCVAHIGPVDSGNHNLITNNHDQHQRRNNSNSSATPIVQFQGESNCLKCF	300
	RYRLNDRHRHLFDLISSTWHWASSKAPHKHAIVTVTYDSEEQRQQFLDVVKIPPTISHKL	360
	GFMSLHLL 368	
	HPV6B E4	
30	1 MGAPNIGKYVMAAQLYVLLHLYLALHKKYPFLNLLHTPPHRPPPLCPQAPRKTQCKRRLG	60
	NEHEESNSPLATPCVWPTLDPWTVETTTSSLTITTSTKDGTTVTVQLRL 109	
	HPV6B ESA	
	1 MEVVPVQIAAGTTSTFILPVIIAFVVCFVSIILIVWISEFIVYTSVLVLTLLLYLLLWLL	60
35	LTTPLQFFLLTLLVCYCPALYIHYYIVTTQQ 91	
	HPV6B E5B	
	1 MMLTCQFNDGDTWLGLWLLCAFIVGMLGLLLMHYRAVQGDKHTKCKKCNKHNCNDDYVTM	60
	HYTTDGDYIYMN 72	
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	HPV6B E6	
	1 MESANASTSATTIDQLCKTFNLSMHTLQINCVFCKNALTTAEIYSYAYKHLKVLFRGGYP	60
	YAACACCLEFHGKINQYRHFDYAGYATTVEEETKQDILDVLIRCYLCHKPLCEVEKVKHI	120
	LTKARFIKLNCTWKGRCLHCWTTCMEDMLP 150	
5		
	HPV6B E7	
	1 MHGRHVTLKDIVLDLQPPDPVGLHCYEQLVDSSEDEVDEVDGQDSQPLKQHFQIVTCCCG	60
	CDSNVRLVVQCTETDIREVQQLLLGTLNIVCPICAPKT 98	
10	HPV6A L1	
	1 MWRPSDSTVYVPPPNPVSKVVATDAYVTRTNIFYHASSSRLLAVGHPYFSIKRANKTVVP	60
	KVSGYQYRVFKVVLPDPNKFALPDSSLFDPTTQRLVWACTGLEVGRGQPLGVGVSGHPFL	120
	NKYDDVENSGSGGNPGQDNRVNVGMDYKQTQLCMVGCAPPLGEHWGKGKQCTNTPVQAGD	180
	CPPLELITSVIQDGDMVDTGFGAMNFADLQTNKSDVPIDICGTTCKYPDYLQMAADPYGD	240
15	RLFFFLRKEQMFARHFFNRAGEVGEPVPDTLIIKGSGNRTSVGSSIYVNTPSGSLVSSEA	300
	QLFNKPYWLQKAQGHNNGICWGNQLFVTVVDTTRSTNMTLCASVTTSSTYTNSDYKEYMR	360
	HVEEYDLQFIFQLCSITLSAEVMAYIHTMNPSVLEDWNFGLSPPPNGTLEDTYRYVQSQA	420
	ITCQKPTPEKEKPDPYKNLSFWEVNLKEKFSSELDQYPLGRKFLLQSGYRGRSSIRTGVK	480
	RPAVSKASAAPKRKRAKTKR 500	
20		
	HPV6B L2	
	1 MAHSRARRKRASATQLYQTCKLTGTCPPDVIPKVEHNTIADQILKWGSLGVFFGGLGIG	60
	TGSGTGGRTGYVPLQTSAKPSITSGPMARPPVVVEPVAPSDPSIVSLIEESAIINAGAPE	120
	IVPPAHGGFTITSSETTTPAILDVSVTSHTTTSIFRNPVFTEPSVTQPQPPVEANGHILI	180
25	SAPTVTSHPIEEIPLDTFVVSSSDSGPTSSTPVPGTAPRPRVGLYSRALHQVQVTDPAFL	240
	STPORLITYDNPVYEGEDVSVQFSHDSIHNAPDEAFMDIIRLHRPAIASRRGLVRYSRIG	300
	QRGSMHTRSGKHIGARIHYFYDISPIAQAAEEIEMHPLVAAQDDTFDIYAESFEPGINPT	360
	OHPVTNISDTYLTSTPNTVTQPWGNTTVPLSLPNDLFLQSGPDITFPTAPMGTPFSPVTP	420
	ALPTGPVFITGSGFYLHPAWYFARKRRKRIPLFFSDVAA 453	
30		
	HPV11 E1	
	1 MADDSGTENEGSGCTGWFMVEAIVEHTTGTQISEDEEEEVEDSGYDMVDFIDDRHITQNS	60
	VEAQALFNRQEADAHYATVQDLKRKYLGSPYVSPISNVANAVESEISPRLDAIKLTTQPK	120
	KVKRRLFETRELTDSGYGYSEVEAATQVEKHGDPENGGDGQERDTGRDIEGEGVEHREAE	180
35	AVDDSTREHADTSGILELLKCKDIRSTLHGKFKDCFGLSFVDLIRPFKSDRTTCADWVVA	240
	GFGIHHSIADAFQKLIEPLSLYAHIQWLTNAWGMVLLVLIRFKVNKSRCTVARTLGTLLN	300
	IPENHMLIEPPKIQSGVRALYWFRTGISNASTVIGEAPEWITRQTVIEHSLADSQFKLTE	360
	MVQWAYDNDICEESEIAFEYAQRGDFDSNARAFLNSNMQAKYVKDCAIMCRHYKHAEMKK	420
	MSIKQWIKYRGTKVDSVGNWKPIVQFLRHQNIEFIPFLSKLKLWLHGTPKKNCIAIVGPP	480
40	DTGKSCFCMSLIKFLGGTVISYVNSCSHFWLQPLTDAKVALLDDATQPCWTYMDTYMRNL	540
	LDGNPMSIDRKHRALTLIKCPPLLVTSNIDISKEEKYKYLHSRVTTFTFPNPFPFDRNGN	600

AVYELSDANWKCFFERLSSSLDIEDSEDEEDGSNSQAFRCVPGSVVRTL 649

	HPV11 E2	
	1 MEAIAKRLDACQDQLLELYEENSIDIHKHIMHWKCIRLESVLLHKAKQMGLSHIGLQVVP	60
5	PLTVSETKGHNAIEMQMHLESLAKTQYGVEPWTLQDTSYEMWLTPPKRCFKKQGNTVEVK	120
	FDGCEDNVMEYVVWTHIYLQDNDSWVKVTSSVDAKGIYYTCGQFKTYYVNFNKEAQKYGS	180
	TNHWEVCYGSTVICSPASVSSTVREVSIAEPTTYTPAQTTAPTVSACTTEDGVSAPPRKR	240
	ARGPSTNNTLCVANIRSVDSTINNIVTDNYNKHQRRNNCHSAATPIVQLQGDSNCLKCFR	300
	YRLNDKYKHLFELASSTWHWASPEAPHKNAIVTLTYSSEEQRQQFLNSVKIPPTIRHKVG	360
10	FMSLHLL 367	
	HPV11 E4	
	1 MVVPIIGKYVMAAQLYVLLHLYLALYEKYPLLNLLHTPPHRPPPLQCPPAPRKTACRRRL	60
	GSEHVDRPLTTPCVWPTSDPWTVQSTTSSLTITTSTKEGTTVTVQLRL 108	
15		
	HPV11 E5A	
	1 MEVVPVQIAAATTTTLILPVVIAFAVCILSIVLIILISDFVVYTSVLVLTLLLYLLLWLL	60
	LTTPLQFFLLTLCVCYFPAFYIHIYIVQTQQ 91	
20	HPV11 E5B	
	${\tt 1} {\tt MVMLTCHLNDGDTWLFLWLFTAFVVAVLGLLLLHYRAVHGTEKTKCAKCKSNRNTTVDYV}$	60
	YMSHGDNGDYVYMN 74	
	HPV11 E6	
25	1 MESKDASTSATSIDQLCKTFNLSLHTLQIQCVFCRNALTTAEIYAYAYKNLKVVWRDNFP	60
	FAACACCLELQGKINQYRHFNYAAYAPTVEEETNEDILKVLIRCYLCHKPLCEIEKLKHI	120
	LGKARFIKLNNQWKGRCLHCWTTCMEDLLP 150	
	HPV11 E7	
30	1 MHGRLVTLKDIVLDLQPPDPVGLHCYEQLEDSSEDEVDKVDKQDAQPLTQHYQILTCCCG	60
	CDSNVRLVVECTDGDIRQLQDLLLGTLNIVCPICAPKP 98	
	HPV11 L1	
	1 MWRPSDSTVYVPPPNPVSKVVATDAYVKRTNIFYHASSSRLLAVGHPYYSIKKVNKTVVP	60
35	${\tt KVSGYQYRVFKVVLPDPNKFALPDSSLFDPTTQRLVWACTGLEVGRGQPLGVGVSGHPLL}$	120
	NKYDDVENSGGYGGNPGQDNRVNVGMDYKQTQLCMVGCAPPLGEHWGKGTQCSNTSVQNG	180
	${\tt DCPPLELITSVIQDGDMVDTGFGAMNFADLQTNKSDVPLDICGTVCKYPDYLQMAADPYG}$	240
	DRLFFYLRKEQMFARHFFNRAGTVGEPVPDDLLVKGGNNRSSVASSIYVHTPSGSLVSSE	300
	${\tt AQLFNKPYWLQKAQGHNNGICWGNHLFVTVVDTTRSTNMTLCASVSKSATYTNSDYKEYM}$	360
40	${\tt RHVEEFDLQFIFQLCSITLSAEVMAYIHTMNPSVLEDWNFGLSPPPNGTLEDTYRYVQSQ}$	420
	A THE OWNER PROPERTY OF THE PR	480

KRPAVSKPSTAPKRKRTKTKK 501

	HPV11 L2	
	1 MKPRARRKRASATQLYQTCKATGTCPPDVIPKVEHTTIADQILKWGSLGVFFGGLGIGT	60
5	GAGSGGRAGYIPLGSSPKPAITGGPAARPPVLVEPVAPSDPSIVSLIEESAIINAGAPEV	120
	VPPTQGGFTITSSESTTPAILDVSVTNHTTTSVFQNPLFTEPSVIQPQPPVEASGHILIS	180
	APTITSQHVEDIPLDTFVVSSSDSGPTSSTPLPRAFPRPRVGLYSRALQQVQVTDPAFLS	240
	TPQRLVTYDNPVYEGEDVSLQFTHESIHNAPDEAFMDIIRLHRPAITSRRGLVRFSRIGQ	300
	RGSMYTRSGQHIGARIHYFQDISPVTQAAEEIELHPLVAAENDTFDIYAEPFDPIPDPVQ	360
0	HSVTQSYLTSTPNTLSQSWGNTTVPLSIPSDWFVQSGPDITFPTASMGTPFSPVTPALPT	420
	GPVFITGSDFYLHPTWYFARRRRKRIPLFFTDVAA 455	
	HPV16 E1	
	1 MADPAGTNGEEGTGCNGWFYVEAVVEKKTGDAISDDENENDSDTGEDLVDFIVNDNDYLT	60
15	QAETETAHALFTAQEAKQHRDAVQVLKRKYLVSPLSDISGCVDNNISPRLKAICIEKQSR	120
	AAKRRLFESEDSGYGNTEVETQQMLQVEGRHETETPCSQYSGGSGGGCSQYSSGSGGEGV	180
	SERHTICQTPLTNILNVLKTSNAKAAMLAKFKELYGVSFSELVRPFKSNKSTCCDWCIAA	240
	FGLTPSIADSIKTLLQQYCLYLHIQSLACSWGMVVLLLVRYKCGKNRETIEKLLSKLLCV	300
	SPMCMMIEPPKLRSTAAALYWYKTGISNISEVYGDTPEWIQRQTVLQHSFNDCTFELSQM	360
20	VQWAYDNDIVDDSEIAYKYAQLADTNSNASAFLKSNSQAKIVKDCATMCRHYKRAEKKQM	420
	SMSQWIKYRCDRVDDGGDWKQIVMFLRYQGVEFMSFLTALKRFLQGIPKKNCILLYGAAN	480
	TGKSLFGMSLMKFLQGSVICFVNSKSHFWLQPLADAKIGMLDDATVPCWNYIDDNLRNAL	540
	DGNLVSMDVKHRPLVQLKCPPLLITSNINAGTDSRWPYLHNRLVVFTFPNEFPFDENGNP	600
	VYELNDKNWKSFFSRTWSRLSLHEDEDKENDGDSLPTFKCVSGQNTNTL 649	
25		
	HPV16 E2 Accession number W2WLHS	
	1 METLCQRLNVCQDKILTHYENDSTDLRDHIDYWKHMRLECAIYYKAREMGFKHINHQVVP	60
	TLAVSKNKALQAIELQLTLETIYNSQYSNEKWTLQDVSLEVYLTAPTGCIKKHGYTVEVQ	120
	FDGDICNTMHYTNWTHIYICEEASVTVVEGQVDYYGLYYVHEGIRTYFVQFKDDAEKYSK	180
30	NKVWEVHAGGQVILCPTSVFSSNEVSSPEIIRQHLANHPAATHTKAVALGTEETQTTIQR	240
	PRSEPDTGNPCHTTKLLHRDSVDSAPILTAFNSSHKGRINCNSNTTPIVHLKGDANTLKC	300
	LRYRFKKHCTLYTAVSSTWHWTGHNVKHKSAIVTLTYDSEWQRDQFLSQVKIPKTITVST	360
	GFMSI 365	
35	HPV16 E5 Accession number W5WLHS	
	1 MTNLDTASTTLLACFLLCFCVLLCVCLLIRPLLLSVSTYTSLIILVLLLWITAASAFRCF	61
	IVYIIFVYIPLFLIHTHARFLIT 83	
	HPV16 E6	

YRDGNPYAVCDKCLKFYSKISEYRHYCYSLYGTTLEQQYNKPLCDLLIRCINCQKPLCPE 120

1 MHQKRTAMFQDPQERPRKLPQLCTELQTTIHDIILECVYCKQQLLRREVYDFAFRDLCIV 60

40

EKQRHLDKKQRFHNIRGRWTGRCMSCCRSSRTRRETQL 158

5	HPV16 E7 1 MHGDTPTLHEYMLDLQPETTDLYCYEQLNDSSEEEDBIDGPAGQAEPDRAHYNIVTFCCK CDSTLRLCVQSTHVDIRTLEDLLMGTLGIVCPICSQKP 98 .	60
10	HPV16 L1 ACCESSION NUMBER AAD33259 1 MQVTFIYILVITCYENDVNVYHIFFQMSLWLPSEATVYLPPVPVSKVVSTDEYVARTNIY YHAGTSRLLAVGHPYFPIKEPINNKILVPKVSGLGYRVFRIHLPPDPKFGFPDTSFYYPD TQRLVWACVGVEVGRGQPLGVGISGHPLLNKLDDTENASAYAANAGVDNRECISMDYKQT QLCLIGCKPPIGEHMGKGSECTNVAVNPGDCPPLELINTVIQDGDWYDDTGFGAMDFTTLQ ANKSEVPLDICTSICKYPDYIKMVSEPYGDSLFFYLRREQMFVRHLFNRAGAVGENVPDD LYIKGSGSTANLASSNYFPTPSGSMVTSDAQIFNKEYMLQRAQGHNWGICMGNQLFVTVV DTTRSTNMSLCAALSTSETTYKNTNFKEYLRHGBEYDLQFIFQLCKITLTADVMTYIHSM NSTILEDWNFGLQPPGGTLEDTYRFVTSQAIACQKHTPPAPKEDPLKKYTFMEVNLKEK FSADLDQFFLGRFFLLQAGLKAKFKFTLGKRKATPTTSSTSTTAKRKKRKL 531	60 120 180 240 300 360 420
20	HPV16 L2 Accession number AAD33258 1 MRHKRSAKRTKRASATQLYKTCKQAGTCPPDIIPKVEKKTIADQILQYGSMGVFFGGLGI GTGGGTGGRTGYIPLGTRPPTATDTLAPVRPPLTVDPVGPSDPSIVSLVEETSFIDAGAP TSVPSIPPDVSGFSITTSTDTTPAILDINNTVTTVTTHNNPTFDPSVLQPPTPAETGGH FTLSSSTISTHNYESIPPDTFIVSTNPNTVTSSTPIPGSRPVARLGLYSRTTQQVKVVDP AFITTPTKLITYDNPAYEGIDVDNTLYFSSNDNSINIAPDPDFLDIVALHRPALTSRTG IRYSRIGNKQTLRTRSGKSIGAKVHYYYDFSTIDSAEEISLQTITPSTYTTTSHAALPTS INNGLYDIYADDFITDTSTTPVPSVPSTELSGYIPANTTIPFGGAYNIPLVSGPDIPINI TDQAPSLIPIVPGSPQYTIIADAGDFYLHPSYYMLRKRKKRLPYFFSDVSLAA 473	60 120 180 240 300 360 420
30	HPV18 E1 1 MADPEGTDGBGTGCNGWFYVQAIVDKKTGDVISDDEDENATDTGSDMVDFIDTQGTFCEQ AELETAQALFHAQEVHNDAQVLHVLKRKEAGGSTENSFLCERLEVDTELSFRLQBISLNS GQKKAKRRLFTISDSGYGCSEVEATQIQVTTNGEHGGNVCSGGSTEAIDNGGTEGNNSSV DGTSDNSNIENVNPQCTIAQLKDLLKVNKQGAMLAVFKDTYGLSFTDLVRNFKSDKTC TDWVTAIFGVMPTIABGFKTLLQFFILVAHIQCLDCKWGVLILALLRYKCGKSRLTVAKG LSTLLHVPETCMLIQPFKLRSSVAALYWYRTGISNISEVMGDTPEWIQRLTIIQHGIDDS NFDLSEMVQWAPDNSLTDESDMAFFYALLADSNSNAAAFLKSNCQAKYLKDCATMCKHYR RAQKRQMMMSQWIRFRCSKIDESGDWRPIVQFLRYQQIEFITFLGALKSFLKGTFKNCL VFCGPANTGKSYFGMSFIHFIQGAVISFVNSTSHFWLEPLTDTKVAMLDDATTTCMTYFD TYMRNALDGNFISIDRKHKFLIQLKCPFILLTINHPAKDNKWPYLSSKITVFFFPNAFP	60 120 180 240 300 360 420 480 540
40	FDKNGNPVYEINDKNWKCFFERTWSRLDLHEEBEDADTEGNPFGTFKLRAGQNHRPL 65	7

	HPV18 E2 Accession number W2WL18	
	1 MQTPKETLSERLSCVQDKIIDHYENDSKDIDSQIQYWQLIRWENAIFFAAREHGIQTLNH	60
	QVVPAYNISKSKAHKAIELQMALQGLAQSRYKTEDWTLQDTCEELWNTEPTHCFKKGGQT	120
	VQVYFDGNKDNCMTYVAWDSVYYMTDAGTWDKTATCVSHRGLYYVKEGYNTFYIEFKSEC	180
5	EKYGNTGTWEVHFGNNVIDCNDSMCSTSDDTVSATQLVKQLQHTPSPYSSTVSVGTAKTY	240
	GQTSAATRPGHCGLAEKQHCGPVNPLLGAATPTGNNKRRKLCSGNTTPIIHLKGDRNSLK	300
	CLRYRLRKHSDHYRDISSTWHWTGAGNEKTGILTVTYHSETQRTKFLNTVAIPDSVQILV	360
	GYMTM 365	
10	HPV18 E5 Accession number W5WL18	
	1 MLSLIFLFCFCVCMYVCCHVPLLPSVCMCAYAWVLVFVYIVVITSPATAFTVYVFCFLLP	60
	MLLLHIHAILSLQ 73	
	HPV18 E6	
15	1 MARFEDPTRRPYKLPDLCTELNTSLQDIEITCVYCKTVLELTEVFEFAFKDLFVVYRDSI	60
	PHAACHKCIDFYSRIRELRHYSDSVYGDTLEKLTNTGLYNLLIRCLRCQKPLNPAEKLRH	120
	LNEKRRFHNIAGHYRGQCHSCCNRARQERLQRRRETQV 158	
	HPV18 E7	
20	1 MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT	60
	MLCMCCKCEARIKLVVESSADDLRAFQQLFLNTLSFVCPWCASQQ 105	
	HPV18 L1 Accession number CAA28671	
	1 MCLYTRVLILHYHLLPLYGPLYHPRPLPLHSILVYMVHIIICGHYIILFLRNVNVFPIFL	60
25	QMALWRPSDNTVYLPPPSVARVVNTDDYVTPTSIFYHAGSSRLLTVGNPYFRVPAGGGNK	120
	QDIPKVSAYQYRVFRVQLPDPNKFGLPDTSIYNPETQRLVWACAGVEIGRGQPLGVGLSG	180
	HPFYNKLDDTESSHAATSNVSEDVRDNVSVDYKQTQLCILGCAPAIGEHWAKGTACKSRP	240
	LSQGDCPPLELKNTVLEDGDMVDTGYGAMDFSTLQDTKCEVPLDICQSICKYPDYLQMSA	300
	DPYGDSMFFCLRREQLFARHFWNRAGTMGDTVPQSLYIKGTGMPASPGSCVYSPSPSGSI	360
30	VTSDSQLFNKPYWLHKAQGHNNGVCWHNQLFVTVVDTTPSTNLTICASTQSPVPGQYDAT	420
	KFKQYSRHVEEYDLQFIFQLCTITLTADVMSYIHSMNSSILEDWNFGVPPPPTTSLVDTY	480
	RFVQSVAITCQKDAAPAENKDPYDKLKFWNVDLKEKFSLDLDQYPLGRKFLVQAGLRRKP	540
	TIGPRKRSAPSATTSSKPAKRVRVRARK 568	
35	HPV18 L2 Accession number P2WL18	
	1 MVSHRAARRKRASVTDLYKTCKQSGTCPPDVVPKVEGTTLADKILQWSSLGIFLGGLGIG	60
	TGSGTGGRTGYIPLGGRSNTVVDVGPTRPPVVIEPVGPTDPSIVTLIEDSSVVTSGAPRP	120
	TFTGTSGFDITSAGTTTPAVLDITPSSTSVSISTTNFTNPAFSDPSIIEVPQTGEVAGNV	180
	FVGTPTSGTHGYEEIPLQTFASSGTGEEPISSTPLPTVRRVAGPRLYSRAYQQVSVANPE	240
40	${\tt FLTRPSSLITYDNPAFEPVDTTLTFDPRSDVPDSDFMDIIRLHRPALTSRRGTVRFSRLG}$	300
	ORATMFTRSGTQIGARVHFYHDISPIAPSPEYIELQPLVSATEDNDLFDIYADDMDPAVP	360

	VPSRSTTSFAFFKYSPTISSASSYSNVTVPLTSSWDVPVYTGPDITLPSTTSVWPIVSPT	420
	APASTQYIGIHGTHYYLWPLYYFIPKKRKRVPYFFADGFVAA 462	
	HPV31 E1 Accession number W1WL31	
5	1 MADPAGTDGEGTGCNGWFYVEAVIDRQTGDNISEDENEDSSDTGEDMVDFIDNCNVYNNQ	60
	AEAETAQALFHAQEAEEHAEAVQVLKRKYVGSPLSDISSCVDYNISPRLKAICIENNSKT	120
	AKRRLFELPDSGYGNTEVETQQMVQVEEQQTTLSCNGSDGTHSERENETPTRNILQVLKT	180
	SNGKAAMLGKFKELYGVSFMELIRPFQSNKSTCTDWCVAAFGVTGTVAEGFKTLLQPYCL	240
	YCHLQSLACSWGMVMLMLVRFKCAKNRITIEKLLEKLLCISTNCMLIQPPKLRSTAAALY	300
10	WYRTGMSNISDVYGETPEWIERQTVLQHSFNDTTFDLSQMVQWAYDNDVMDDSEIAYKYA	360
	QLADSDSNACAFLKSNSQAKIVKDCGTMCRHYKRAEKRQMSMGQWIKSRCDKVSDEGDWR	420
	DIVKFLRYQQIEFVSFLSALKLFLKGVPKKNCILIHGAPNTGKSYFGMSLISFLQGCIIS	480
	YANSKSHFWLQPLADAKIGMLDDATTPCWHYIDNYLRNALDGNPVSIDVKHKALMQLKCP	540
	PLLITSNINAGKDDRWPYLHSRLVVFTFPNPFPFDKNGNPVYELSDKNWKSFFSRTWCRL	600
15	NLHEEEDKENDGDSFSTFKCVSGQNIRTL 629	
	HPV31 E2 Accession number W2WL31	
	1 METLSQRLNVCQDKILEHYENDSKRLCDHIDYWKHIRLECVLMYKAREMGIHSINHQVVP	60
	ALSVSKAKALQAIELQMMLETLNNTEYKNEDWTMQQTSLELYLTAPTGCLKKHGYTVEVQ	120
20	FDGDVHNTMHYTNWKFIYLCIDGQCTVVEGQVNCKGIYYVHEGHITYFVNFTEEAKKYGT	180
	GKKWEVHAGGQVIVFPESVFSSDEISFAGIVTKLPTANNTTTSNSKTCALGTSEGVRRAT	240
	TSTKRPRTEPEHRNTHHPNKLLRGDSVDSVNCGVISAAACTNQTRAVSCPATTPIIHLKG	300
	DANILKCLRYRLSKYKQLYEQVSSTWHWTCTDGKHKNAIVTLTYISTSQRDDFLNTVKIP	360
	NTVSVSTGYMTI 372	
25		
	HPV31 E5 Accession number W5WL31	
	1 MIELNISTVSIVLCFLLCFCVLLFVCLVIRPLVLSVSVYATLLLLIVILWVIATSPLRCF	60
	CIYVVFIYIPLFVIHTHASFLSQQ 84	
30	HPV31 E6 Accession number W6WL31	
	1 MFKNPAERPRKLHELSSALEIPYDELRLNCVYCKGQLTETEVLDFAFTDLTIVYRDDTPH	60
	GVCTKCLRFYSKVSEFRWYRYSVYGTTLEKLTNKGICDLLIRCITCQRPLCPEEKQRHLD	120
	KKKRFHNIGGRWTGRCIACWRRPRTETQV 149	
35	HPV31 E7 Accession number W7WL31	
	1 MRGETPTLQDYVLDLQPEATDLHCYEQLPDSSDEEDVIDSPAGQAEPDTSNYNIVTFCCQ	60
	CKSTLRLCVQSTQVDIRILQELLMGSFGIVCPNCSTRL 98	
	HPV31 L1 Accession number P1WL31	
40	1 MSLWRPSEATVYLPPVPVSKVVSTDEYVTRTNIYYHAGSARLLTVGHPYYSIPKSDNPKK	60
. •	IVVPKVSGLQYRVFRVRLPDPNKFGFPDTSFYNPETQRLVWACVGLEVGRGQPLGVGISG	120

	HPLLNKFDDTENSNRYAGGPGTDNRECISMDYKQTQLCLLGCKPPIGEHWGKGSPCSNNA	180
	ITPGDCPPLELKNSVIQDGDMVDTGFGAMDFTALQDTKSNVPLDICNSICKYPDYLKMVA	240
	EPYGDTLFFYLRREQMFVRHFFNRSGTVGESVPTDLY1KGSGSTATLANSTYFPTPSGSM	300
	VTSDAQIFNKPYWMQRAQGHNNGICWGNQLFVTVVDTTRSTNMSVCAAIANSDTTFKSSN	360
5	FKEYLRHGEEFDLQFIFQLCKITLSADIMTYIHSMNPAILEDWNFGLTTPPSGSLEDTYR	420
	PVTSQAITCQKTAPQKPKEDPFKDYVFWEVNLKEKFSADLDQFPLGRKFLLQAGYRARPK	480
	FKAGKRSAPSASTTTPAKRKKTKK 504	
	HPV31 L2 Accession number P2WL31	
10	1 MRSKRSTKRTKRASATQLYQTCKAAGTCPSDVIPKIEHTTIADQILRYGSMGVFFGGLGI	60
	GSGSGTGGRTGYVPLSTRPSTVSEASIPIRPPVSIDPVGPLDPSIVSLVEESGIVDVGAP	120
	APIPHPPTTSGFDIATTADTTPAILDVTSVSTHENPTFTDPSVLQPPTPAETSGHLLLSS	180
	SSISTHNYEEIPMDTFIVSTNNENITSSTPIPGVRRPARLGLYSKATQQVKVIDPTFLSA	240
	PKQLITYENPAYETVNAEESLYFSNTSHNIAPDPDFLDIIALHRPALTSRRNTVRYSRLG	300
15	NKQTLRTRSGATIGARVHYYYDISSINPAGESIEMQPLGASATTTSTLNDGLYDIYADTD	360
	FTVDTPATHNVSPSTAVQSTSAVSAYVPTNTTVPLSTGFDIPIFSGPDVPIEHAPTQVFP	420
	FPLAPTTPQVSIFVDGGDFYLHPSYYMLKRRRKRVSYFFTDVSVAA 466	
	HPV45 E1 Accession number S36563	
20	1 MADPEGTDGEGTGCNGWFFVETIVEKKTGDVISDDEDETATDTGSDMVDFIDTQLSICEQ	60
	AEQETAQALFHAQEVQNDAQVLHLLKRKFAGGSKENSPLGEQLSVDTDLSPRLQEISLNS	120
	GHKKAKRRLFTISDSGYGCSEVEAAETQVTVNTNAENGGSVHSTQSSGGDSSDNAENVDP	180
	HCSITELKELLQASNKKAAMLAVFKDIYGLSFTDLVRNFKSDKTTCTDWVMAIFGVNPTV	240
	AEGFKTLIKPATLYAHIQCLDCKWGVLILALLRYKCGKNRLTVAKGLSTLLHVPETCMLI	300
25	EPPKLRSSVAALYWYRTGISNISEVSGDTPEWIQRLTIIQHGIDDSNFDLSDMVQWAFDN	360
	DLTDESDMAFQYAQLADCNSNAAAFLKSNCQAKYLKDCAVMCRHYKRAQKRQMNMSQWIK	420
	YRCSKIDEGGDWRPIVQFLRYQGVEFISFLRALKEFLKGTPKKNCILLYGPANTGKSYFG	480
	MSFIHFLQGAIISFVNSNSHFWLEPLADTKVAMLDDATHTCWTYFDNYMRNALDGNPISI	540
	DRKHKPLLQLKCPPILLTSNIDPAKDNKWPYLESRVTVFTFPHAFPFDKNGNPVYEINDK	600
30	NWKCFFERTWSRLDLHEDDEDADTEGIPFGTFKCVTGQNTRPL 643	
	HPV45 E2 Accession number S36564	
	MKMQTPKESLSERLSALQDKILDHYENDSKDINSQISYWQLIRLENAILFTAREHGITKL	60
	${\tt NHQVVPPINISKSKAHKAIELQMALKGLAQSKYNNEEWTLQDTCEELWNTEPSQCFKKGG}$	120
35	KTVHVYFDGNKDNCMNYVVWDSIYYITETGIWDKTAACVSYWGVYYIKDGDTTYYVQFKS	180
	ECEKYGNSNTWEVQYGGNVIDCNDSMCSTSDDTVSATQIVRQLQHASTSTPKTASVGTPK	240
	PHIQTPATKRPRQCGLTEQHHGRVNTHVHNPLLCSSTSNNKRRKVCSGNTTPIIHLKGDK	300
	$\tt NSLKCLRYRLRKYADHYSEISSTWHWTGCNKNTGLLTVTYNSEVQRNTFLDVVTIPNSVQ$	360
	ISVGYMTI 368	
40		

	HPV45 E6 Accession number CAB44706	
	1 MARFDDPTQRPYKLPDLCTELNTSLQDVSIACVYCKATLERTEVYQFAFKDLFIVYRDCI	60
	AYAACHKCIDFYSRIRELRYYSNSVYGETLEKITNTELYNLLIRCLRCQKPLNPAEKRRH	120
	LKDKRRFHS1AGQYRGQCNTCCDQARQERLRRRRETQV 158	
5		
	HPV45 E7 Accession number CAB44707	
	1 MHGPRATLQEIVLHLEPQNELDPVDLLCYEQLSESEEENDEADGVSHAQLPARRAEPQRH	60
	KILCVCCKCDGRIELTVESSADDLRTLQQLFLSTLSFVCPWCATNQ 106	
10	HPV45 L1 Accession number CAB44705	
	1 MAHNIIYGHGIIIFLKNVNVFPIFLQMALWRPSDSTVYLPPPSVARVVNTDDYVSRTSIF	60
	YHAGSSRLLTVGNPYFRVVPSGAGNKQAVPKVSAYQYRVFRVALPDPNKFGLPDSTIYNP	120
	ETQRLVWACVGMEIGRGQPLGIGLSGHPFYNKLDDTESAHAATAVITQDVRDNVSVDYKQ	180
	TQLCILGCVPAIGEHWAKGTLCKPAQLQPGDCPPLELKNTIIEDGDMVDTGYGAMDFSTL	240
15	QDTKCEVPLD1CQS1CKYPDYLQMSADPYGDSMFFCLRREQLFARHFWNRAGVMGDTVPT	300
	DLYIKGTSANMRETPGSCVYSPSPSGSITTSDSQLFNKPYWLHKAQGHNNGICWHNQLFV	360
	TVVDTTRSTNLTLCASTQNPVPNTYDPTKFKHYSRHVEEYDLQFIFQLCTITLTAEVMSY	420
	IHSMNSSILENWNFGVPPPPTTSLVDTYRFVQSVAVTCQKDTTPPEKQDPYDKLKFWTVD	480
	LKEKFSSDLDQYPLGRKFLVQAGLRRRPTIGPRKRPAASTSTASRPAKRVRIRSKK 536	
20		
	HPV45 L2 Accession number S36565	
	1 MVSHRAARRKRASATDLYRTCKQSGTCPPDVINKVEGTTLADKILQWSSLGIFLGGLGIG	60
	TGSGSGGRTGYVPLGGRSNTVVDVGPTRPPVVIEPVGPTDPSIVTLVEDSSVVASGAPVP	120
	TFTGTSGFEITSSGTTTPAVLDITPTVDSVSISSTSFTNPAFSDPSIIEVPQTGEVSGNI	180
25	FVGTPTSGSHGYEEIPLQTFASSGSGTEPISSTPLPTVRRVRGPRLYSRANQQVRVSTSQ	240
	FLTHPSSLVTFDNPAYEPLDTTLSFEPTSNVPDSDFMDIIRLHRPALSSRRGTVRFSRLG	300
	QRATMFTRSGKQIGGRVHFYHDISPIAATEEIELQPLISATNDSDLFDVYADFPPPASTT	360
	PSTIHKSFTYPKYSLTMPSTAASSYSNVTVPLTSAWDVPIYTGPDIILPSHTPMWPSTSP	420
	TNASTTTYIGIHGTQYYLWPWYYYFPKKRKRIPYFFADGFVAA 463	
30		
	HPV33 El Accession number W1WL33	
	1 MADPEGTNGAGMGCTGWFEVEAVIERRTGDNISEDEDETADDSGTDLLEFIDDSMENSIQ	60
	ADTEAARALFNIQEGEDDLNAVCALKRKFAACSQSAAEDVVDRAANPCRTSINKNKECTY	120
	RKRKIDELEDSGYGNTEVETQQMVQQVESQNGDTNLNDLESSGVGDDSEVSCETNVDSCE	180
35	NVTLQEISNVLHSSNTKANILYKFKEAYGISFMELVRPFKSDKTSCTDWCITGYGISPSV	240
	AESLKVLIKQHSLYTHLQCLTCDRGIIILLLIRFRCSKNRLTVAKLMSNLLSIPETCMVI	300
	EPPKLRSQTCALYWFRTAMSNISDVQGTTPEWIDRLTVLQHSFNDNIFDLSEMVQWAYDN	360
	ELTDDSDIAYYYAQLADSNSNAAAFLKSNSQAKIVKDCGIMCRHYKKAEKRKMSIGQWIQ	420
	SRCEKTNDGGNWRPIVQLLRYQNIEFTAFLGAFKKFLKGIPKKSCMLICGPANTGKSYFG	480
40	MSLIQFLKGCVISCVNSKSHFWLQPLSDAKIGMIDDVTPISWTYIDDYMRNALDGNEISI	540
	DVKHRALVQLKCPPLLLTSNTNAGTDSRWPYLHSRLTVFEFKNPFPFDENGNPVYAINDE	600

NWKSFFSRTWCKLDLIEEEDKENHGGNISTFKCSAGENTRSLRS 644

	HPV33 E2 Accession number W2WL33	
	1 MEEISARLNAVQEKILDLYEADKTDLPSQIEHWKLIRMECALLYTAKQMGFSHLCHQVVP	60
5	SLLASKTKAFQVIELQMALETLSKSQYSTSQWTLQQTSLEVWLCEPPKCFKKQGETVTVQ	120
	YDNDKKNTMDYTNWGEIYIIEEDTCTMVTGKVDYIGMYYIHNCEKVYFKYFKEDAAKYSK	180
	TQMWEVHVGGQVIVCPTSISSNQISTTETADIQTDNDNRPPQAAAKRRRPADTTDTAQPL	240
	TKLFCADPALDNRTARTATNCTNKQRTVCSSNVAPIVHLKGESNSLKCLRYRLKPYKELY	300
	SSMSSTWHWTSDNKNSKNGIVTVTFVTEQQQQMFLGTVKIPPTVQISTGFMTL 353	
10	•	
	HPV33 E5 Accession number W5WL33	
	1 MIFVFVLCFILFLCLSLLLRPLILSISTYAWLLVLVLLLWVFVGSPLKIFFCYLLFLYLP	60
	MMCINFHAQHMTQQE 75	
15	HPV33 E6 Accession number W6WL33	
	${\tt 1} {\tt MFQDTEEKPRTLHDLCQALETTIHNIELQCVECKKPLQRSEVYDFAFADLTVVYREGNPF}$	60
	GICKLCLRFLSKISEYRHYNYSVYGNTLEQTVKKPLNEILIRCIICQRPLCPQEKKRHVD	120
	LNKRFHNISGRWAGRCAACWRSRRRETAL 149	
20	HPV33 E7 Accession number W7WL33	
	1 MRGHKPTLKEYVLDLYPEPTDLYCYEQLSDSSDEDEGLDRPDGQAQPATADYYIVTCCHT	60
	CNTTVRLCVNSTASDLRTIQQLLMGTVNIVCPTCAQQ 97	
	HPV33 L1 Accession number P1WL33	60
25	1 MSVWRPSEATVYLPPVPVSKVVSTDEYVSRTSIYYYAGSSRLLAVGHPYFSIKNPTNAKK	120
	LLVPKVSGLQYRVFRVRLPDPNKFGFPDTSFYNPDTQRLVWACVGLEIGRGQPLGVGISG	180
	HPLLNKFDDTETGNKYPGQPGADNRECLSMDYKQTQLCLLGCKPPTGEHWGKGVACTNAA	240
	PANDCPPLELINTIIEDGDMVDTGFGCMDFKTLQANKSDVPIDICGSTCKYPDYLKMTSE	300
	PYGDSLFFFLRREQMFVRHFFNRAGTLGEAVPDDLYIKGSGTTASIQSSAFFPTPSGSMV	360
30	TSESQLFNKPYWLQRAQGHNNGICWGNQVFVTVVDTTRSTNMTLCTQVTSDSTYKNENFK	420
	EYIRHVEEYDLQFVFQLCKVTLTAEVMTYIHAMNPDILEDWQFGLTPPPSASLQDTYRFV	480
	TSQAITCQKTVPPKEKEDPLGKYTFWEVDLKEKFSADLDQFPLGRKFLLQAGLKAKPKLK	400
	RAAPTSTRTSSAKRKKVKK 499	
	Term 2.2	
35	HPV33 L2 Accession number P2WL33	60
	1 MRHKRSTRRKRASATQLYQTCKATGTCPPDVIPKVEGSTIADQILKYGSLGVFFGGLGIG	120
	TGSGSGGRTGYVPIGTDPPTAAIPLQPIRPPVTVDTVGPLDSSIVSLIEETSFIEAGAPA	180
	PS1PTPSGFDVTTSADTTPAIINVSSVGESSIQTISTHLNPTFTEPSVLHPPAPAEASGH FIFSSPTVSTQSYENIPMDTFVVSTDSSNVTSSTPIPGSRPVARLGLYSRNTQQVKVVDP	240
40	FIFSSPTVSTQSYENIPMDTFVVSTDSSNVTSSTPIFGSRPVARLGGISRATQQVXVVDF AFLTSPHKLITYDNPAFESFDPEDTLQFQHSDISPAPDPDFLDIIALHRPAITSRRHTVR	300
40	AFLTSPHKLITYDNPAFESFDPEDTLQFQASDISFAFDPDFEDITALAKFATISKANIVA FSRVGQKATLKTRSGKQIGARIHYYQDLSPIVPLDHTVPNEQYELQPLHDTSTSSYSIND	360
	FSRVGQKATLKTKSGKQIGAKIHIIQDDSPIVPDDHIVPNEQIEDQFHMDISISSISIND	550

	GLYDVYADDVDNVHTPMQHSYSTFATTRTSNVSIPLNTGFDTPVMSGPDIPSPLFPTSSP	420
	FVPISPFFPFDTIVVDGADFVLHPSYFILRRRKRFPYFFTDVRVAA 467	
	HPV56 E2 Accession number S36581	
5	1 MVPCLOVCKAKACSAIEVQIALESLSTTIYNNEEWTLRDTCEELWLTEPKKCFKKEGQHI	60
,	EVWFDGSKNNCMQYVAWKYIYYNGDCGWQKVCSGVDYRGIYYVHDGHKTYYTDFEQEAKK	120
	FGCKNIWEVHMENESIYCPDSVSSTCRYNVSPVETVNEYNTHKTTTTTSTSVGNQDAAVS	180
	HRPGKRPRLRESEFDSSRESHAKCVTTHTHISDTDNTDSRSRSINNNNHPGDKTTPVVHL	240
	KGEPNRLKCCRYRFQKYKTLFVDVTSTYHWTSTDNKNYSIITIIYKDETQRNSFLSHVKI	300
10	PVVYRLVWDK 310	
10	POVIREMENT	
	HPV56 E6 Accession number W6WL56	
	1 MEPQFNNPQERPRSLHHLSEVLEIPLIDLRLSCVYCKKELTRAEVYNFACTELKLVYRDD	60
	FPYAVCRVCLLFYSKVRKYRYYDYSVYGATLESITKKQLCDLLIRCYRCQSPLTPEEKQL	120
15	HCDRKRRFHLIAHGWTGSCLGCWRQTSREPRESTV 155	
	C36500	
	HPV56 E7 Accession number S36580 1 MHGKVPTLQDVVLELTPQTEIDLQCNEQLDSSEDEDEDEVDHLQERPQQARQAKQHTCYL	60
	IHVPCCECKFVVQLDIQSTKEDLRVVQQLLMGALTVTCPLCASSN 105	
20		
	HPV56 L1 Accession number S38563	60
	1 MMLPMMYIYRDPPLHYGLCIFLDVGAVNVFPIFLQMATWRPSENKVYLPPTPVSKVVATD	120
	SYVKRTSIFYHAGSSRLLAVGHPYYSVTKDNTKTNIPKVSAYQYRVFRVRLPDPNKFGLP	180
	DTNIYNPDQERLVWACVGLEVGRGQPLGAGLSGHPLFNRLDDTESSNLANNNVIEDSRDN	240
25	ISVDGKQTQLCIVGCTPAMGEHWTKGAVCKSTQVTTGDCPPLALINTPIEDGDMIDTGFG	
	AMDFKVLQESKAEVPLDIVQSTCKYPDYLKMSADAYGDSMWFYLRREQLFARHYFNRAGK	300
	VGETIPAELYLKGSNGREPPPSSVYVATPSGSMITSEAQLFNKPYWLQRAQGHNNGICWG	360
	NQLFVTVVDTTRSTNMTISTATEQLSKYDARKINQYLRHVEEYELQFVFQLCKITLSAEV	420
	MAYLHNMNANLLEDWNIGLSPPVATSLEDKYRYVRSTAITCQREQPPTEKQDPLAKYKFW	480
30	DVNLQDSFSTDLDQFPLGRKFLMQLGTRSKPAVATSKKRSAPTSTSTPAKRKRR 534	
	HPV56 L2 Accession number S36582	
	1 MVAHRATRRKRASATQLYKTCKLSGTCPEDVVNKIEQKTWADKILQWGSLFTYFGGLGIG	60
	TGTGSGGRAGYVPLGSRPSTIVDVTPARPPIVVESVGPTDPSIVTLVEESSVIESGAGIP	120
35	NFTGSGGFEITSSSTTTPAVLDITPTSSTVHVSSTHITNPLFIDPPVIEAPQTGEVSGNI	180
55	LISTPTSGIHSYEEIPMQTFAVHGSGTEPISSTPIPGFRRIAAPRLYRKAFQQVKVTDPA	240
	FLDRPATLVSADNPLFEGTDTSLAFSPSGVAPDPDFMNIVALHRPAFTTRRGGVRFSRLG	300
	RKATIQTRRGTQIGARVHYYYDISPIAQAEEIEMQPLLSANNSFDGLYDIYANIDDEAPG	360
	LSSQSVATPSAHLPIKPSTLSFASNTTNVTAPLGNVWETPFYSGPDIVLPTGPSTWPFVP	420
40	OSPYDVTHDVYIQGSSFALWPVYFFRRRRKIPYFFADGDVAA 464	
40	ÖSSIDAIUDAII ÖGGOLUDUL ATI INGGOGGGATI II	

HLA Class I Motifs Indicative of CTL Inducing Peptide Epitopes:

The primary anchor residues of the HLA class I peptide epitope supermotifs and motifs delineated below are summarized in Table I. The HLA class I motifs set out in Table I(a) are those most particularly relevant to the invention claimed here. Primary and secondary anchor positions are summarized in Table II. Allele-specific HLA molecules that comprise HLA class I supertype families are listed in Table VI. In some cases, peptide epitopes may be listed in both a motif and a supermotif Table. The relationship of a particular motif and respective supermotif is indicated in the description of the individual motifs.

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III.D.1. HLA-A1 supermotif The HLA-A1 supermotif is characterized by the presence in peptide ligands of a small (T or S) or hydrophobic (L, I, V, or M) primary anchor residue in position 2, and an aromatic (Y, F, or W) primary anchor residue at the C-terminal position of the epitope. The corresponding family of HLA molecules that bind to the A1 supermotif (i.e., the HLA-A1 supertype) is comprised of at least A*0101, A*2601, A*2602, A*2501, and A*3201 (see, e.g., DiBrino, M. et al., J. Immunol. 151:5930, 1993; DiBrino, M. et al., J. Immunol. 152:620, 1994; Kondo, A. et al., Immunogenetics 45:249, 1997). Other allele-specific HLA molecules predicted to be members of the A1 superfamily are shown in Table VI. Peptides binding to each of the individual HLA proteins can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif. Representative peptide epitopes that comprise the A1 supermotif are set forth in Table VII.

III.D.2. HLA-A2 supermotif

Primary anchor specificities for allele-specific HLA-A2.1 molecules (see, e.g., Falk et al., Nature 351:290-296, 1991; Hunt et al., Science 255:1261-1263, 1992; Parker et al., J. Immunol. 149:3580-25 3587, 1992; Ruppert et al., Cell 74:929-937, 1993) and cross-reactive binding among HLA-A2 and -A28 molecules have been described. (See, e.g., Fruci et al., Human Immunol. 38:187-192, 1993; Tanigaki et al., Human Immunol. 39:155-162, 1994; Del Guercio et al., J. Immunol. 154:685-693, 1995; Kast et al., J. Immunol. 152:3904-3912, 1994 for reviews of relevant data.) These primary anchor residues define the HLA-A2 supermotif; which presence in peptide ligands corresponds to the ability to bind several different 30 HLA-A2 and -A28 molecules. The HLA-A2 supermotif comprises peptide ligands with L, I, V, M, A, T, or Q as a primary anchor residue at position 2 and L, I, V, M, A, or T as a primary anchor residue at the Cterminal position of the epitope.

The corresponding family of HLA molecules (i.e., the HLA-A2 supertype that binds these peptides) is comprised of at least: A*0201, A*0202, A*0203, A*0204, A*0205, A*0206, A*0207, A*0209, A*0214, A*6802, and A*6901. Other allele-specific HLA molecules predicted to be members of the A2 superfamily are shown in Table VI. As explained in detail below, binding to each of the individual allele-specific HLA molecules can be modulated by substitutions at the primary anchor and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise an A2 supermotif are set forth in Table VIII. The motifs comprising the primary anchor residues V, A, T, or Q at position 2 and L, I, V, A, or T at the C-terminal position are those most particularly relevant to the invention claimed herein.

5 III.D.3. HLA-A3 supermotif

The HLA-A3 supermotif is characterized by the presence in peptide ligands of A, L, I, V, M, S, or, T as a primary anchor at position 2, and a positively charged residue, R or K, at the C-terminal position of the epitope, e.g., in position 9 of 9-mers (see, e.g., Sidney et al., Hum. Immunol. 45:79, 1996). Exemplary members of the corresponding family of HLA molecules (the HLA-A3 supertype) that bind the A3 supermotif include at least A*0301, A*1101, A*3101, A*3301, and A*6801. Other allele-specific HLA molecules predicted to be members of the A3 supertype are shown in Table VI. As explained in detail below, peptide binding to each of the individual allele-specific HLA proteins can be modulated by substitutions of amino acids at the primary and/or secondary anchor positions of the peptide, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise the A3 supermotif are set forth in Table IX.

III.D.4. HLA-A24 supermotif

The HLA-A24 supermotif is characterized by the presence in peptide ligands of an aromatic (F, W, or Y) or hydrophobic allphatic (L, I, V, M, or T) residue as a primary anchor in position 2, and Y, F, W, L, I, or M as primary anchor at the C-terminal position of the epitope (see, e.g., Sette and Sidney, Immunogenetics 1999 Nov;50/3-4):201-12, Review). The corresponding family of HLA molecules that bind to the A24 supermotif (i.e., the A24 supertype) includes at least A*2402, A*3001, and A*2301. Other allele-specific HLA molecules predicted to be members of the A24 supertype are shown in Table VI. Peptide binding to each of the allele-specific HLA molecules can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise the A24 supermotif are set forth in Table

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III.D.5. HLA-B7 supermotif

The HLA-B7 supermotif is characterized by peptides bearing prolline in position 2 as a primary anchor, and a hydrophobic or aliphatic amino acid (L, I, V, M, A, F, W, or Y) as the primary anchor at the C-terminal position of the epitope. The corresponding family of HLA molecules that bind the B7 supermotif (i.e., the HLA-B7 supertype) is comprised of a least twenty six HLA-B proteins including: B+0702, B+0703, B+0704, B+1508, B+3501, B+3502, B*3503, B*3504, B*3505, B*3506, B*3507, B*3508, B*5101, B*5102, B*5103, B*5104, B*5105, B*5301, B*5401, B*5501, B*5502, B*5601, B*5601, B*5701, and B*7801 (see, e.g., Sidney, et al., J. Immunol. 154:247, 1995; Barber, et al., Curr. Biol. 5:179, 1995; Hill, et al., Nature 360:434, 1992; Rammensee, et al., Immunogenetics 41:178, 1995 for reviews of relevant data). Other allele-specific HLA molecules predicted to be members of the B7 supertype are shown in Table VI. As explained in detail below, peptide binding to each of the individual

allele-specific HLA proteins can be modulated by substitutions at the primary and/or secondary anchor positions of the peptide, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise the B7 supermotif are set forth in Table XI.

5 III.D.6, HLA-B27 supermotif

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The HLA-B27 supermotif is characterized by the presence in peptide ligands of a positively charged (R, H, or K) residue as a primary anchor at position 2, and a hydrophobic (F, Y, L, W, M, I, A, or V) residue as a primary anchor at the C-terminal position of the epitope (see, e.g., Sidney and Sette, Immunogenetics 1999 Nov;50(3-4):201-12, Review). Exemplary members of the corresponding family of HLA molecules that bind to the B27 supermoif (t.e., the B27 supertype) include at least B*1401, B*1402, B*1509, B*2702, B*2703, B*2704, B*2705, B*2706, B*3801, B*3901, B*3902, and B*7301. Other allele-specific HLA molecules predicted to be members of the B27 supertype are shown in Table VI. Peptide binding to each of the allele-specific HLA molecules can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative peptide entitopes that comprise the B27 supermotif are set forth in Table

III.D.7. HLA-B44 supermotif

The HLA-B44 supermotif is characterized by the presence in peptide ligands of negatively charged (D or E) residues as a primary anchor in position 2, and hydrophobic residues (F, W, Y, L, I, M, V, or A) as a primary anchor at the C-terminal position of the epitope (see, e.g., Sidney et al., Immunol. Today 17:261, 1996). Exemplary members of the corresponding family of HLA molecules that bind to the B44 supermotif (i.e., the B44 supertype) include at least: B*1801, B*1802, B*3701, B*4001, B*4002, B*4403, and B*4006. Peptide binding to each of the allele-specific HLA molecules can be modulated by substitutions at primary and/or secondary anchor positions; preferably choosing respective residues specified for the supermotif.

III.D.8. HLA-B58 supermotif

The HLA-B58 supermotif is characterized by the presence in peptide ligands of a small aliphatic residue (A, S, or T) as a primary anchor residue at position 2, and an aromatic or hydrophobic residue (F, W, Y, L, I, V, M, or A) as a primary anchor residue at the C-terminal position of the epitope (see, e.g., Sidney and Sette, Immunogenetics 1999 Nov;50(3-4):201-12, Review). Exemplary members of the corresponding family of HLA molecules that bind to the B58 supermotif (i.e., the B58 supertype) include at least B*1516, B*1517, B*5701, B*5702, and B*5801. Other allele-specific HLA molecules predicted to be members of the B58 supertype are shown in Table VI. Peptide binding to each of the allele-specific HLA molecules can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise the B58 supermotif are set forth in Table

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III.D.9. HLA-B62 supermotif

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The HLA-B62 supermotif is characterized by the presence in peptide ligands of the polar aliphatic residue Q or a hydrophobic aliphatic residue (L, V, M, I, or P) as a primary anchor in position 2, and a hydrophobic residue (F, W, Y, M, I, V, L, or A) as a primary anchor at the C-terminal position of the epitope (see, e.g., Sidney and Sette, Immunogenetics 1999 Nov. 50(3-4):201-12. Review). Exemplary members of the corresponding family of HLA molecules that bind to the B62 supermotif (i.e., the B62 supertype) include at least: B*1501, B*1502, B*1513, and B5201. Other allele-specific HLA molecules predicted to be members of the B62 supertype are shown in Table VI. Peptide binding to each of the allele-specific HLA molecules can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative peptide epitopes that comprise the B62 supermotif are set forth in Table XIV.

III.D.10. HLA-A1 motif

The HLA-A1 motif is characterized by the presence in peptide ligands of T, S, or M as a primary anchor residue at position 2 and the presence of Y as a primary anchor residue at the C-terminal position of the epitope. An alternative allele-specific A1 motif is characterized by a primary anchor residue at position 3 rather than position 2. This motif is characterized by the presence of D, E, A, or S as a primary anchor residue in position 3, and a Y as a primary anchor residue at the C-terminal position of the epitope (see, e.g., DiBrino et al., J. Immunol.; 152:620, 1994; Kondo et al., Immunogenetics 45:249, 1997; and Kubo et al., J. Immunol. 152:3913, 1994 for reviews of relevant data). Peptide binding to HLA A1 can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the motif.

Representative peptide epitopes that comprise either A1 motif are set forth in Table XV.

Those epitopes comprising T, S, or M at position 2 and Y at the C-terminal position are also included in the listing of HLA-A1 supermotif-bearing peptides listed in Table VII, as these residues are a subset of the A1 supermotif primary anchors.

III.D.11. HLA-A*0201 motif

An HLA-A2*0201 motif was determined to be characterized by the presence in peptide ligands of L or M as a primary anchor residue in position 2, and L or V as a primary anchor residue at the C-terminal position of a 9-residue peptide (see, e.g., Falk et al., Nature 351:290-296, 1991) and was further found to comprise an 1 at position 2 and I or A at the C-terminal position of a nine amino acid peptide (see, e.g., Hunt et al., Science 255:1261-1263, March 6, 1992; Parker et al., J. Immunol. 149:3580-3587, 1992). The A*0201 allele-specific motif has also been defined by the present inventors to additionally comprise V, A, T, or Q as a primary anchor residue at the C-terminal position of the epitope (see, e.g., Kast et al., J. Immunol. 152:3904-3912, 1994). Thus, the HLA-A*0201 motif comprises peptide ligands with L, I, V, M, A, T, or Q as primary anchor residues at position 2 and L, I, V, M, A, or T as a primary anchor residue at the C-terminal position of the epitope. The preferred and tolerated residues that characterize the primary anchor positions of the HLA-A*0201 motif

are identical to the residues describing the A2 supermotif. (For reviews of relevant data, see, e.g., Del Guercio et al., J. Immunol. 154:683-693, 1995; Ruppert et al., Cell 74:929-937, 1993; Sidney et al., Immunol. 70day 17:261-266, 1996; Sette and Sidney, Curr. Opin. in Immunol. 10:478-482, 1998). Secondary anchor residues that characterize the A*0201 motif have additionally been defined (see, e.g., Ruppert et al., Cell 74:929-937, 1993). These are shown in Table II. Peptide binding to HLA-A*0201 molecules can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the motif.

Representative peptide epitopes that comprise an A*0201 motif are set forth in Table

VIII. The A*0201 motifs comprising the primary anchor residues V, A, T, or Q at position 2 and L, I, V, A,

or T at the C-terminal position are those most particularly relevant to the invention claimed herein.

III.D.12. HLA-A3 motif

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The HLA-A3 motif is characterized by the presence in peptide ligands of L, M, V, I, S, A,
T, F, C, G, or D as a primary anchor residue at position 2, and the presence of K, Y, R, H, F, or A as a
15 primary anchor residue at the C-terminal position of the epitope (see, e.g., DiBrino et al., Proc. Natl. Acad.
Sci USA 90:1508, 1993; and Kubo et al., J. Immunol. 152:3913-3924, 1994). Peptide binding to HLA-A3
can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing
respective residues specified for the motif.

Representative peptide epitopes that comprise the A3 motif are set forth in Table XVI.

Those epitopes that also comprise the A3 supermotif are also listed in Table IX. The A3 supermotif primary anchor residues comprise a subset of the A3- and A11-allele specific motif primary anchor residues.

III.D.13. HLA-A11 motif

The HLA-A11 motif is characterized by the presence in peptide ligands of V, T, M, L, I, S, A, G, N, C, D, or F as a primary anchor residue in position 2, and K, R, Y, or H as a primary anchor residue at the C-terminal position of the epitope (see, e.g., Zhang et al., Proc. Natl. Acad. Sci. USA 90:2217-2221, 1993; and Kubo et al., J. Immunol. 152:3913-3924, 1994). Peptide binding to HLA-A11 can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the motif.

Representative peptide epitopes that comprise the A11 motif are set forth in Table XVII; peptide epitopes comprising the A3 allele-specific motif are also present in this Table because of the extensive overlap between the A3 and A11 motif primary anchor specificities. Further, those peptide epitopes that comprise the A3 supermotif are also listed in Table IX.

III.D.14. HLA-A24 motif

The HLA-A24 motif is characterized by the presence in peptide ligands of Y, F, W, or M as a primary anchor residue in position 2, and F, L, I, or W as a primary anchor residue at the C-terminal position of the epitope (see, e.g., Kondo et al., J. Immunol. 155:4307-4312, 1995; and Kubo et al., J. Immunol. 152:3913-3924, 1994). Peptide binding to HLA-A24 molecules can be modulated by

substitutions at primary and/or secondary anchor positions; preferably choosing respective residues specified for the motif.

Representative peptide epitopes that comprise the A24 motif are set forth in Table XVIII. These epitopes are also listed in Table X, which sets forth HLA-A24-supermotif-bearing peptides, as the primary anchor residues characterizing the A24 allele-specific motif comprise a subset of the A24 supermotif primary anchor residues.

Motifs Indicative of Class II HTL Inducing Peptide Epitopes

The primary and secondary anchor residues of the HLA class Π peptide epitope supermotifs and motifs delineated below are summarized in Table III.

III.D.15. HLA DR-1-4-7 supermotif

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Motifs have also been identified for peptides that bind to three common HLA class II allele-specific HLA molecules: HLA DRB1*0401, DRB1*0101, and DRB1*0701 (see, e.g., the review by Southwood et al. J. Immunology 160:3363-3373,1998). Collectively, the common residues from these motifs delineate the HLA DR-14-7 supermotif. Peptides that bind to these DR molecules carry a supermotif characterized by a large aromatic or hydrophobic residue (Y, F, W, L, I, V, or M) as a primary anchor residue in position 1, and a small, non-charged residue (S, T, C, A, P, V, I, L, or M) as a primary anchor residue in position 6 of a 9-mer core region. Allele-specific secondary effects and secondary anchors for each of these HLA types have also been identified (Southwood et al., supra). These are set forth in Table III. Peptide binding to HLA-DRB1*0401, DRB1*0101, and/or DRB1*0701 can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the supermotif.

Representative 9-mer epitopes comprising the DR-1-4-7 supermotif, wherein position 1 of the supermotif is at position 1 of the nine-residue core, are set forth in Table XIX. Exemplary epitopes of 15 amino acids in length that comprises the nine residue core include the three residues on either side that flank the nine residue core. HTL epitopes that comprise the core sequences can also be of lengths other than 15 amino acids, supra. Accordingly, epitopes of the invention include sequences that typically comprise the nine residue core plus 1, 2, 3 (as in the exemplary 15-mer), 4, or 5 flanking residues on either side of the nine residue core.

III.D.16. HLA DR3 motifs

Two alternative motifs (i.e., submotifs) characterize peptide epitopes that bind to HLADR3 molecules (see, e.g., Geluk et al., J. Immunol. 152:5742, 1994). In the first motif (submotif DR3A) a
35 large, hydrophobic residue (L, 1, V, M, F, or Y) is present in anchor position 1 of a 9-mer core, and D is
present as an anchor at position 4, towards the carboxyl terminus of the epitope. As in other class II motifs,
core position 1 may or may not occupy the peptide N-terminal position.

The alternative DR3 submotti provides for lack of the large, hydrophobic residue at anchor position 1, and/or lack of the negatively charged or amide-like anchor residue at position 4, by the presence of a positive charge at position 6 towards the carboxyl terminus of the epitope. Thus, for the

alternative allele-specific DR3 motif (submotif DR3B): L, I, V, M, F, Y, A, or Y is present at anchor position I; D, N, Q, E, S, or T is present at anchor position 4; and K, R, or H is present at anchor position 6. Peptide binding to HLA-DR3 can be modulated by substitutions at primary and/or secondary anchor positions, preferably choosing respective residues specified for the motif.

Representative 9-mer epitopes corresponding to a nine residue sequence comprising the DR3a and DR3b submotifs (wherein position 1 of the motif is at position 1 of the nine residue core) are set forth in Table XXa and b. Exemplary epitopes of 15 amino acids in length that comprises the nine residue core include the three residues on either side that flank the nine residue core. HTL epitopes that comprises the cores sequences can also be of lengths other than 15 amino acids, supra. Accordingly, epitopes of the invention include sequences that typically comprise the nine residue core plus 1, 2, 3 (as in the exemplary 15-mer), 4, or 5 flanking residues on either side of the nine residue core.

Each of the HLA class I or class II epitopes set out in the Tables herein are deemed singly to be an inventive aspect of this application. Further, it is also an inventive aspect of this application that each epitope may be used in combination with any other epitope.

III.E. Enhancing Population Coverage of the Vaccine

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Vaccines that have broad population coverage are preferred because they are more commercially viable and generally applicable to the most people. Broad population coverage can be obtained using the peptides of the invention (and nucleic acid compositions that encode such peptides) through selecting peptide epitopes that bind to HLA alleles which, when considered in total, are present in most of the population. The Table below lists the overall frequencies of the HLA class I supertypes in various ethnicities (section a) and the combined population coverage achieved by the A2-, A3-, and B7-supertypes (section b). The A2-, A3-, and B7 supertypes are each present on the average of over 40% in each of these five major ethnic groups. Coverage in excess of 80% is achieved with a combination of these supermotifs. These results suggest that effective and non-ethnically biased population coverage is achieved upon use of a limited number of cross-reactive peptides. Although the population coverage reached with these three main peptide specificities is high, coverage can be expanded to reach 95% population coverage and above, and more easily achieve truly multispecific responses upon use of additional supermotif or allele-specific motif bearing peptides.

The B44-, A1-, and A24-supertypes are each present, on average, in a range from 25% to 40% in these major ethnic populations (section a). While less prevalent overall, the B27-, B58-, and B62 supertypes are each present with a frequency >25% in at least one major ethnic group (section a). In section b, the Table summarizes the estimated prevalence of combinations of HLA supertypes that have been identified in five major ethnic groups. The incremental coverage obtained by the inclusion of A1-, A24-, and B44-supertypes to the A2, A3, and B7 coverage and coverage obtained with all of the supertypes described herein, is shown.

The data presented herein, together with the previous definition of the A2-, A3-, and B7supertypes, indicates that all antigens, with the possible exception of A29, B8, and B46, can be classified
into a total of nine HLA supertypes. By including epitopes from the six most frequent supertypes, an
average population coverage of 99% is obtained for five major ethnic groups.

Population coverage with combined HLA Supertypes

	PHENOTYPIC FREQUENCY					
	Caucasian	North	Japanese	Chinese	Hispanic	Average
HLA-SUPERTYPES		American				
		Black				
a. Individual Supertypes						
A2	45.8	39.0	42.4	45.9 ·	43.0	43.2
A3	37.5	42.1	45.8	52.7	43.1	44.2
B7	43.2	55.1	57.1	43.0	49.3	49.5
A1	47.1	16.1	21.8	14.7	26.3	25.2
A24	23.9	38.9	58.6	40.1	38.3	40.0
B44	43.0	21.2	42.9	39.1	39.0	37.0
B27	28.4	26.1	13.3	. 13.9	35.3	23.4
B62	12.6	4.8	36.5	25.4	11.1	18.1
B58	10.0	25.1	1.6	9.0	5.9	10.3
b. Combined Supertypes						
A2, A3, B7	84.3	86.8	89.5	89.8	86.8	87.4
A2, A3, B7, A24, B44, A1	99.5	98.1	100.0	99.5	99.4	99.3
A2, A3, B7, A24, B44, A1, B27, B62, B58	99.9	99.6	100.0	99.8	99.9	99.8

III.F. Immune Response-Stimulating Peptide Analogs

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In general, CTL and HTL responses to whole antigens are not directed against all possible epitopes. Rather, they are restricted to a few "immunodominant" determinants (Zinkermagel, et al., Adv. Immunol. 27:5159, 1979; Bennink, et al., J. Exp. Med. 168:19351939, 1988; Rawle, et al., J. Immunol. 146:397-3984, 1991). It has been recognized that immunodominance (Benacerraf, et al., Science 175:273-279, 1972) could be explained by either the ability of a given epitope to selectively bind a particular HLA protein (determinant selection theory) (Vitello, et al., J. Immunol. 131:1635, 1983); Rosenthal, et al., Nature 267:156-158, 1977), or to be selectively recognized by the existing TCR (T cell receptor) specificities (repertoire theory) (Klein, I., IMMUNOLOGY, THE SCIENCE OF SELENONSELF DISCRIMINATION, John Wiley & Sons, New York, pp. 270-310, 1982). It has been demonstrated that additional factors, mostly linked to processing events, can also play a key role in dictating, beyond strict immunogenicity, which of the many potential determinants will be presented as immunodominant (Sercarz, et al., Annu. Rev. Immunol. 11:729-766, 1993).

The concept of dominance and subdominance is relevant to immunotherapy of both infectious diseases and cancer. For example, in the course of chronic viral disease, recruitment of subdominant epitopes can be important for successful clearance of the infection, especially if dominant CTL or HTL specificities have been inactivated by functional tolerance, suppression, mutation of viruses and other mechanisms (Franco, et al., Curr. Opin. Immunol. 7:524-531, 1995). In the case of cancer and tumor antigens, CTLs recognizing at least some of the highest binding affinity peptides might be functionally inactivated. Lower binding affinity peptides are preferrentially recognized at these times, and may therefore be preferred in therapeutic or prophylactic anti-cancer vaccines.

25 In particular, it has been noted that a significant number of epitopes derived from known non-viral tumor associated antigens (TAA) bind HLA class I with intermediate affinity (IC₃₀ in the 50-500

nM range). For example, it has been found that 8 of 15 known TAA peptides recognized by tumor infiltrating lymphocytes (TIL) or CTL bound in the 50-500 nM range. (These data are in contrast with estimates that 90% of known viral antigens were bound by HLA class I molecules with IC50 of 50 nM or less, while only approximately 10% bound in the 50-500 nM range (Sette, et al., J. Immunol., 153:558-5592, 1994). In the cancer setting this phenomenon is probably due to elimination or functional inhibition of the CTL recognizing several of the highest binding peptides, presumably because of T cell tolerization events.

Without intending to be bound by theory, it is believed that because T cells to dominant epitopes may have been clonally deleted, selecting subdominant epitopes may allow existing T cells to be recruited, which will then lead to a therapeutic or prophylactic response. However, the binding of HLA 10 molecules to subdominant epitopes is often less vigorous than to dominant ones. Accordingly, there is a need to be able to modulate the binding affinity of particular immunogenic epitopes for one or more HLA molecules, and thereby to modulate the immune response elicited by the peptide, for example to prepare analog peptides which elicit a more vigorous response. This ability would greatly enhance the usefulness of peptide epitope-based vaccines and therapeutic agents.

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Although peptides with suitable cross-reactivity among all alleles of a superfamily are identified by the screening procedures described above, cross-reactivity is not always as complete as possible, and in certain cases procedures to increase cross-reactivity of peptides can be useful; moreover, such procedures can also be used to modify other properties of the peptides such as binding affinity or peptide stability. Having established the general rules that govern cross-reactivity of peptides for HLA alleles within a given motif or supermotif, modification (i.e., analoging) of the structure of peptides of particular interest in order to achieve broader (or otherwise modified) HLA binding capacity can be performed. More specifically, peptides which exhibit the broadest cross-reactivity patterns, can be produced in accordance with the teachings herein. The present concepts related to analog generation are set forth in greater detail in co-pending U.S.S.N. 09/226,775 filed 1/6/99.

In brief, the strategy employed utilizes the motifs or supermotifs which correlate with binding to certain HLA molecules. The motifs or supermotifs are defined by having primary anchors, and in many cases secondary anchors. Analog peptides can be created by substituting amino acid residues at primary anchor, secondary anchor, or at primary and secondary anchor positions. Generally, analogs are made for peptides that already bear a motif or supermotif. Preferred secondary anchor residues of supermotifs and motifs that have been defined for HLA class I and class II binding peptides are shown in Tables II and III, respectively.

For a number of the motifs or supermotifs in accordance with the invention, residues are defined which are deleterious to binding to allele-specific HLA molecules or members of HLA supertypes that bind the respective motif or supermotif (Tables II and III). Accordingly, removal of such residues that are detrimental to binding can be performed in accordance with the present invention. For example, in the case of the A3 supertype, when all peptides that have such deleterious residues are removed from the population of peptides used in the analysis, the incidence of cross-reactivity increased from 22% to 37% (see, e.g., Sidney, J. et al., Hu. Immunol. 45:79, 1996). Thus, one strategy to improve the cross-reactivity of peptides within a given supermotif is simply to delete one or more of the deleterious residues present

within a peptide and substitute a small "neutral" residue such as Ala (that may not influence T cell recognition of the peptide). An enhanced likelihood of cross-reactivity is expected if, together with elimination of detrumental residues within a peptide, "preferred" residues associated with high affinity binding to an allele-specific HLA molecule or to multiple HLA molecules within a superfamily are inserted.

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To ensure that an analog peptide, when used as a vaccine, actually elicits a CTL response to the native epitope in vivo (or, in the case of class II epitopes, elicits helper T cells that cross-react with the wild type peptides), the analog peptide may be used to immunize T cells in vitro from individuals of the appropriate HLA allele. Thereafter, the immunized cells' capacity to induce lysis of wild type peptide sensitized target cells is evaluated. It will be desirable to use as antigen presenting cells, cells that have been either infected, or transfected with the appropriate genes, or, in the case of class II epitopes only, cells that have been pulsed with whole protein antigens, to establish whether endogenously produced antigen is also recognized by the relevant T cells.

Another embodiment of the invention is to create analogs of weak binding peptides, to thereby ensure adequate numbers of cross-reactive cellular binders. Class I binding peptides exhibiting binding affinities of 500-5000 nM, and carrying an acceptable but suboptimal primary anchor residue at one or both positions can be "fixed" by substituting preferred anchor residues in accordance with the respective supertype. The analog peptides can then be tested for crossbinding activity.

Another embodiment for generating effective peptide analogs involves the substitution of residues that have an adverse impact on peptide stability or solubility in, e.g., a liquid environment. This substitution may occur at any position of the peptide epitope. For example, a cysteine (C) can be substituted out in favor of \(\alpha \)-amino butyric acid. Due to its chemical nature, cysteine has the propensity to form disulfide bridges and sufficiently alter the peptide structurally so as to reduce binding capacity. Substituting \(\alpha \)-amino butyric acid for \(\Cappa \) not only alleviates this problem, but actually improves binding and crossbinding capability in certain instances (\(\text{see}, e.g. \), the review by Sette et al., \(\text{In Persistent Yiral Infections} \). Eds. R. Ahmed and I. Chen, John Wiley & Sons, England, 1999). Substitution of cysteine with \(\alpha \)-amino butyric acid may occur at any residue of a peptide epitope, i.e. at either anchor or non-anchor positions.

III.G. Computer Screening of Protein Sequences from Disease-Related Antigens for Supermotif- or 30 Motif-Bearing Peptides

In order to identify supermotif- or motif-bearing epitopes in a target antigen, a native protein sequence, e.g., a tumor-associated antigen, or sequences from an infectious organism, or a donor tissue for transplantation, is screened using a means for computing, such as an intellectual calculation or a computer, to determine the presence of a supermotif or motif within the sequence. The information obtained from the analysis of native peptide can be used directly to evaluate the status of the native peptide or may be utilized subsequently to generate the peptide epitope.

Computer programs that allow the rapid screening of protein sequences for the occurrence of the subject supermotifs or motifs are encompassed by the present invention; as are programs that permit the generation of analog peptides. These programs are implemented to analyze any identified amino acid sequence or operate on an unknown sequence and simultaneously determine the sequence and identify

motif-bearing epitopes thereof; analogs can be simultaneously determined as well. Generally, the identified sequences will be from a pathogenic organism or a tumor-associated peptide. For example, the target molecules considered herein include, without limitation, the E1, E2, E4, E5a, E5b, E6, E7, L1 and L2 proteins of HPV.

5 In cases where the sequences of multiple variants of the same target protein are available, potential peptide epitopes can also be selected on the basis of their conservancy. For example, a criterion for conservancy may define that the entire sequence of an HLA class I binding peptide or the entire 9-mer core of a class II binding peptide, be conserved in a designated percentage, of the sequences evaluated for a specific protein antigen.

To target a broad population that may be infected with a number of different strains, it is preferable to include in vaccine compositions epitopes that are representative of HPV antigen sequences from different HPV strains. As appreciated by those in the art, regions with greater or lessor degrees of conservancy among HPv strains can be employed as appropriate for a given antigenic target.

It is important that the selection criteria utilized for prediction of peptide binding are as

accurate as possible, to correlate most efficiently with actual binding. Prediction of peptides that bind, for
example, to HLA-A*0201, on the basis of the presence of the appropriate primary anchors, is positive at
about a 30% rate (see, e.g., Ruppert, J. et al. Cell 74:929, 1993). However, by extensively analyzing
peptide-HLA binding data disclosed herein, data in related patent applications, and data in the art, the
present inventors have developed a number of allele-specific polynomial algorithms that dramatically
increase the predictive value over identification on the basis of the presence of primary anchor residues
alone. These algorithms take into account not only the presence or absence of primary anchors, but also
consider the positive or deleterious presence of secondary anchor residues (to account for the impact of
different amino acids at different positions). The algorithms are essentially based on the premise that the
overall affinity (or \(\Delta \text{O} \) of peptide-HLA interactions can be approximated as a linear polynomial function of

 $\Delta G = a_{ii} \times a_{2i} \times a_{3i} ... \times a_{ni}$

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where a_{ij} is a coefficient that represents the effect of the presence of a given amino acid (i) at a given position (i) along the sequence of a peptide of n amino acids. An important assumption of this method is that the effects at each position are essentially independent of each other. This assumption is justified by studies that demonstrated that peptides are bound to HLA molecules and recognized by T cells in essentially an extended conformation. Derivation of specific algorithm coefficients has been described, for example, in Gulukota, K. et al., J. Mol. Biol. 267:1258, 1997.

Additional methods to identify preferred peptide sequences, which also make use of specific motifs, include the use of neural networks and molecular modeling programs (see, e.g., Milik et al., Nature Biotechnology 16:753, 1998; Altuvia et al., Hum. Immunol. 58:1, 1997; Altuvia et al., J. Mol. Biol. 249:244, 1995; Buus, S. Curr. Opin. Immunol. 11:209-213, 1999; Brusic, V. et al., Bioinformatics 14:121-130, 1998; Parker et al., J. Immunol. 152:163, 1993; Meister et al., Vaccine 13:581, 1995; Hammer et al., J. Exp. Med. 180:2353, 1994; Sturniolo et al., Nature Biotechnol. 17:555 1999).

For example, it has been shown that in sets of A*0201 motif-bearing peptides containing
40 at least one preferred secondary anchor residue while avoiding the presence of any deleterious secondary

anchor residues, 69% of the peptides will bind A $^{\circ}$ 0201 with an IC₃₀ less than 500 nM (Ruppert, J. et al. Cell 74:929, 1993). These algorithms are also flexible in that cut-off scores may be adjusted to select sets of peptides with greater or lower predicted binding properties, as desired.

In utilizing computer screening to identify peptide epitopes, a protein sequence or

stranslated sequence may be analyzed using software developed to search for motifs, for example the

"FINDPATTERNS' program (Devereux, et al. Nucl. Acids Res. 12:387-395, 1984) or MotifSearch 1.4

software program (D. Brown, San Diego, CA) to identify potential peptide sequences containing
appropriate HLA binding motifs. The identified peptides can be scored using customized polynomial
algorithms to predict their capacity to bind specific HLA class I or class II alleles. As appreciated by one of
ordinary skill in the art, a large array of computer programming software and hardware options are available
in the relevant art which can be employed to implement the motifs of the invention in order to evaluate
(e.g., without limitation, to identify epitopes, identify epitope concentration per peptide length, or to
generate analogs) known or unknown peptide sequences.

In accordance with the procedures described above, HPV peptide epitopes that are able to bind HLA supertype groups or allele-specific HLA molecules have been identified (Tables VII-XX).

III.H. Preparation of Peptide Epitopes

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Peptides in accordance with the invention can be prepared synthetically, by recombinant DNA technology or chemical synthesis, or from natural sources such as native tumors or pathogenic organisms. Peptide epitopes may be synthesized individually or as polyepitopic peptides. Although the peptide will preferably be substantially free of other naturally occurring host cell proteins and fragments thereof, in some embodiments the peptides may be synthetically conjugated to native fragments or particles.

The petides in accordance with the invention can be a variety of lengths, and either in

their neutral (uncharged) forms or in forms which are salts. The peptides in accordance with the invention are either free of modifications such as glycosylation, side chain oxidation, or phosphorylation; or they contain these modifications, subject to the condition that modifications do not destroy the biological activity of the peptides as described herein.

When possible, it may be desirable to optimize HLA class I binding epitopes of the invention, such as can be used in a polyepitopic construct, to a length of about 8 to about 13 amino acid residues, often 8 to 11, preferably 9 to 10. HLA class II binding peptide epitopes of the invention may be optimized to a length of about 6 to about 30 amino acids in length, preferably to between about 13 and about 20 residues. Preferably, the peptide epitopes are commensurate in size with endogenously processed pathogen-derived peptides or tumor cell peptides that are bound to the relevant HLA molecules, however, the identification and preparation of peptides that comprise epitopes of the invention can also be carried out using the techniques described herein.

In alternative embodiments, epitopes of the invention can be linked as a polyepitopic peptide, or as a minigene that encodes a polyepitopic peptide.

In another embodiment, it is preferred to identify native peptide regions that contain a high concentration of class I and/or class II epitopes. Such a sequence is generally selected on the basis that it contains the greatest number of epitopes per amino acid length. It is to be appreciated that epitopes can

be present in a nested or overlapping manner, e.g. a 10 amino acid long peptide could contain two 9 amino acid long epitopes and one 10 amino acid long pitope; upon intracellular processing, each epitope can be exposed and bound by an HLA molecule upon administration of such a peptide. This larger, preferably multi-epitopic, peptide can be generated synthetically, recombinantly, or via cleavage from the native source.

The peptides of the invention can be prepared in a wide variety of ways. For the preferred relatively short size, the peptides can be synthesized in solution or on a solid support in accordance with conventional techniques. Various automatic synthesizers are commercially available and can be used in accordance with known protocols. (See, for example, Stewart & Young, SOLID PHASE PEPTIDE SYNTHESIS, 2D. ED., Pierce Chemical Co., 1984). Further, individual peptide epitopes can be joined using chemical ligation to produce larger peptides that are still within the bounds of the invention.

Alternatively, recombinant DNA technology can be employed wherein a nucleotide sequence which encodes an immunogenic peptide of interest is inserted into an expression vector, transformed or transfected into an appropriate host cell and cultivated under conditions suitable for expression. These procedures are generally known in the art, as described generally in Sambrook et al., MOLECULAR CLONING, A LABORATORY MANUAL, Cold Spring Harbor Press, Cold Spring Harbor, New York (1989). Thus, recombinant polypeptides which comprise one or more peptide sequences of the invention can be used to present the appropriate T cell epitope.

The nucleotide coding sequence for peptide epitopes of the preferred lengths contemplated herein can be synthesized by chemical techniques, for example, the phosphotriester method of Matteucci, et al., J. Am. Chem. Soc. 103:3185 (1981). Peptide analogs can be made simply by substituting the appropriate and desired nucleic acid base(s) for those that encode the native peptide sequence; exemplary nucleic acid substitutions are those that encode an amino acid defined by the motifs/supermotifs herein. The coding sequence can then be provided with appropriate linkers and ligated into expression vectors commonly available in the art, and the vectors used to transform suitable hosts to produce the desired fusion protein. A number of such vectors and suitable host systems are now available. For expression of the fusion proteins, the coding sequence will be provided with operably linked start and stop codons, promoter and terminator regions and usually a replication system to provide an expression vector for expression in the desired cellular host. For example, promoter sequences compatible with bacterial hosts are provided in plasmids containing convenient restriction sites for insertion of the desired coding sequence. The resulting expression vectors are transformed into suitable bacterial hosts. Of course, yeast, insect or mammalian cell hosts may also be used, employing suitable vectors and control sequences.

III.I. Assays to Detect T-Cell Responses

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Once HLA binding peptides are identified, they can be tested for the ability to elicit a Tcell response. The preparation and evaluation of motif-bearing peptides are described in PCT publications
WO 94/20127 and WO 94/03205. Briefly, peptides comprising epitopes from a particular antigen are
synthesized and tested for their ability to bind to the appropriate HLA proteins. These assays may involve
evaluating the binding of a peptide of the invention to purified HLA class I molecules in relation to the
binding of a radioiodinated reference peptide. Alternatively, cells expressing empty class I molecules (i.e.

lacking peptide therein) may be evaluated for peptide binding by immunofluorescent staining and flow microfluorimetry. Other assays that may be used to evaluate peptide binding include peptide-dependent class I assembly assays and/or the inhibition of CTL recognition by peptide competition. Those peptides that bind to the class I molecule, typically with an affinity of 500 nM or less, are further evaluated for their ability to serve as targets for CTLs derived from infected or immunized individuals, as well as for their capacity to induce primary in vitro or in vivo CTL responses that can give rise to CTL populations capable of reacting with selected target cells associated with a disease.

Analogous assays are used for evaluation of HLA class II binding peptides. HLA class II motif-bearing peptides that are shown to bind, typically at an affinity of 1000 nM or less, are further evaluated for the ability to stimulate HTL responses.

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Conventional assays utilized to detect T cell responses include proliferation assays, lymphokine secretion assays, direct cytotoxicity assays, and limiting dilution assays. For example, antigen-presenting cells that have been incubated with a peptide can be assayed for the ability to induce CTL responses in responder cell populations. Antigen-presenting cells can be normal cells such as peripheral blood mononuclear cells or dendritic cells. Alternatively, mutant non-human mammalian cell lines that are deficient in their ability to load class I molecules with internally processed peptides and that have been transfected with the appropriate human class I gene, may be used to test for the capacity of the peptide to induce in vitro primary CTL responses.

Peripheral blood mononuclear cells (PBMCs) may be used as the responder cell source of

CTL precursors. The appropriate antigen-presenting cells are incubated with peptide, after which the
peptide-loaded antigen-presenting cells are then incubated with the responder cell population under
optimized culture conditions. Positive CTL activation can be determined by assaying the culture for the
presence of CTLs that kill radio-labeled target cells, both specific peptide-pulsed targets as well as target
cells expressing endogenously processed forms of the antigen from which the peptide sequence was

derived.

Additionally, a method has been devised which allows direct quantification of antigenspecific T cells by staining with Fluorescein-labelled HLA tetrameric complexes (Altman, J. D. et al., Proc.
Natl. Acad. Sci. USA 90:10330, 1993; Altman, J. D. et al., Science 274:94, 1996). Other relatively recent
technical developments include staining for intracellular lymphokines, and interferon release assays or
ELISPOT assays. Tetramer staining, intracellular lymphokine staining and ELISPOT assays all appear to
be at least 10-fold more sensitive than more conventional assays (Lalvani, A. et al., J. Exp. Med. 186:859,
1997; Dunbar, P. R. et al., Curr. Biol. 8:413, 1998; Murali-Krishna, K. et al., Immunity 8:177, 1998).

HTL activation may also be assessed using such techniques known to those in the art such as T cell proliferation and secretion of lymphokines, e.g. IL-2 (see, e.g. Alexander et al., Immunity 1:751-761, 1994).

Alternatively, immunization of HLA transgenic mice can be used to determine immunogenicity of peptide epitopes. Several transgenic mouse models including mice with human A2.1, A11 (which can additionally be used to analyze HLÁ-A3 epitopes), and B7 alleles have been characterized and others (e.g., transgenic mice for HLA-A1 and A24) are being developed. HLA-DR1 and HLA-DR3 mouse models have also been developed. Additional transgenic mouse models with other HLA alleles may

be generated as necessary. Mice may be immunized with peptides emulsified in Incomplete Freund's Adjuvant and the resulting T cells tested for their capacity to recognize peptide-pulsed target cells and target cells transfected with appropriate genes. CTL responses may be analyzed using cytotoxicity assays described above. Similarly, HTL responses may be analyzed using such assays as T cell proliferation or secretion of lymphokines.

III.J. Use of Peptide Epitopes as Diagnostic Agents and for Evaluating Immune Responses In one aspect of the invention, HLA class I and class II binding peptides as described

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herein can be used as reagents to evaluate an immune response. The immune response to be evaluated is induced by using as an immunogen any agent that may result in the production of antigen-specific CTLs or HTLs that recognize and bind to the peptide epitope(s) to be employed as the reagent. The peptide reagent need not be used as the immunogen. Assay systems that are used for such an analysis include relatively recent technical developments such as tetramers, staining for intracellular lymphokines and interferon release assays, or ELISPOT assays.

For example, a peptide of the invention is used in a tetramer staining assay to assess peripheral blood mononuclear cells for the presence of antigen-specific CTLs following exposure to a pathogen or immunogen. The HLA-tetrameric complex is used to directly visualize antigen-specific CTLs (see, e., Ogg et al., Science 279:2103-2106, 1998; and Altman et al., Science 174:94-96, 1996) and determine the frequency of the antigen-specific CTL population in a sample of peripheral blood mononuclear cells.

A tetramer reagent using a peptide of the invention is generated as follows: A peptide that binds to an HLA molecule is refolded in the presence of the corresponding HLA heavy chain and \(\textit{\textit{BL}}_2\) microglobulin to generate a trimolecular complex. The complex is biotinylated at the carboxyl terminal end of the heavy chain at a site that was previously engineered into the protein. Tetramer formation is then induced by the addition of streptavidin. By means of fluorescently labeled streptavidin, the tetramer can be used to stain antigen-specific cells. The cells can then be readily identified, for example, by flow cytometry. Such procedures are used for diagnostic or prognostic purposes. Cells identified by the procedure can also be used for therapeutic purposes.

Peptides of the invention are also used as reagents to evaluate immune recall responses. (see, e.g., Bertoni et al., J. Clin. Invest. 100:503-513, 1997 and Penna et al., J. Exp. Med. 174:1565-1570, 1991.) For example, patient PBMC samples from individuals infected with HPV are analyzed for the presence of antigen-specific CTLs or HTLs using specific peptides. A blood sample containing mononuclear cells may be evaluated by cultivating the PBMCs and stimulating the cells with a peptide of the invention. After an appropriate cultivation period, the expanded cell population may be analyzed, for example, for CTL or for HTL activity.

The peptides are also used as reagents to evaluate the efficacy of a vaccine. PBMCs obtained from a patient vaccinated with an immunogen are analyzed using, for example, either of the methods described above. The patient is HLA typed, and peptide epitope reagents that recognize the allelespecific molecules present in that patient are selected for the analysis. The immunogenicity of the vaccine is indicated by the presence of HPV epitope-specific CTLs and/or HTLs in the PBMC sample.

The peptides of the invention are also be used to make antibodies, using techniques well known in the art (see, e.g. CURRENT PROTOCOLS N IMMUNICOCY, Wiley/Greene, NY; and Antibodies A Laboratory Manual Harlow, Harlow and Lane, Cold Sering Harbor Laboratory Press, 1989), which may be useful as reagents to diagnose HPV infection. Such antibodies include those that recognize a peptide in the context of an HLA molecule, i.e., antibodies that bind to a peptide-MHC complex.

III.K. Vaccine Compositions

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Vaccines and methods of preparing vaccines that contain an immunogenically effective amount of one or more peptides as described herein are further embodiments of the invention. Once appropriately immunogenic epitopes have been defined, they can be sorted and delivered by various means, 10 herein referred to as "vaccine" compositions. Such vaccine compositions can include, for example, lipopeptides (e.g., Vitiello, A. et al., J. Clin. Invest. 95:341, 1995), peptide compositions encapsulated in poly(DL-lactide-co-glycolide) ("PLG") microspheres (see, e.g., Eldridge, et al., Molec. Immunol. 28:287-294, 1991: Alonso et al., Vaccine 12:299-306, 1994; Jones et al., Vaccine 13:675-681, 1995), peptide compositions contained in immune stimulating complexes (ISCOMS) (see, e.g., Takahashi et al., Nature 15 344:873-875, 1990; Hu et al., Clin Exp Immunol. 113:235-243, 1998), multiple antigen peptide systems (MAPs) (see e.g., Tam, J. P., Proc. Natl. Acad. Sci. U.S.A. 85:5409-5413, 1988; Tam, J.P., J. Immunol. Methods 196:17-32, 1996), peptides formulated as multivalent peptides; peptides for use in ballistic delivery systems, typically crystallized peptides, viral delivery vectors (Perkus, M. E. et al., In: Concepts in vaccine development, Kaufmann, S. H. E., ed., p. 379, 1996; Chakrabarti, S. et al., Nature 320:535, 1986; 20 Hu, S. L. et al., Nature 320:537, 1986; Kieny, M.-P. et al., AIDS Bio/Technology 4:790, 1986; Top, F. H. et al. J. Infect. Dis. 124:148, 1971; Chanda, P. K. et al., Virology 175:535, 1990), particles of viral or synthetic origin (e.g., Kofler, N. et al., J. Immunol. Methods. 192:25, 1996; Eldridge, J. H. et al., Sem. Hematol. 30:16, 1993; Falo, L. D., Jr. et al., Nature Med. 7:649, 1995), adjuvants (Warren, H. S., Vogel, F. R., and Chedid, L. A. Annu. Rev. Immunol. 4:369, 1986; Gupta, R. K. et al., Vaccine 11:293, 1993). 25 liposomes (Reddy, R. et al., J. Immunol. 148:1585, 1992; Rock, K. L., Immunol. Today 17:131, 1996), or, naked or particle absorbed cDNA (Ulmer, J. B. et al., Science 259:1745, 1993; Robinson, H. L., Hunt, L. A., and Webster, R. G., Vaccine 11:957, 1993; Shiver, J. W. et al., In: Concepts in vaccine development, Kaufmann, S. H. E., ed., p. 423, 1996; Cease, K. B., and Berzofsky, J. A., Annu. Rev. Immunol. 12:923, 1994 and Eldridge, J. H. et al., Sem. Hematol. 30:16, 1993). Toxin-targeted delivery technologies, also 30 known as receptor mediated targeting, such as those of Avant Immunotherapeutics, Inc. (Needham, Massachusetts) may also be used.

Vaccine compositions of the invention include nucleic acid-mediated modalities. DNA or RNA encoding one or more of the peptides of the invention can also be administered to a patient. This approach is described, for instance, in Wolff et. al., Science 247:1465 (1990) as well as U.S. Patent Nos. 5,880,859; 5,889,465 (5,3804.566; 5,739,118; 5,736,524; 5,679,647; WO 98/04720; and in more detail below. Examples of DNA-based delivery technologies include "naked DNA", facilitated (bupivicaine, polymers, peptide-mediated) delivery, cationic lipid complexes, and particle-mediated ("gene gun") or pressure-mediated delivery (see, e.g., U.S. Patent No. 5,922,687).

For therapeutic or prophylactic immunization purposes, the peptides of the invention can be expressed by viral or bacterial vectors. Examples of expression vectors include attenuated viral hosts, such as vaccinia or fowlpox. This approach involves the use of vaccinia virus, for example, as a vector to express nucleotide sequences that encode the peptides of the invention. Upon introduction into an acutely or chronically infected host or into a non-infected host, the recombinant vaccinia virus expresses the immunogenic peptide, and thereby elicits a host CTL and/or HTL response. Vaccinia vectors and methods useful in immunization protocols are described in, e.g., U.S. Patent No. 4,722,848. Another vector is BCG (Bacille Calmette Guerin). BCG vectors are described in Stover et al., Nature 351:456-460 (1991). A wide variety of other vectors useful for therapeutic administration or immunization of the peptides of the invention, e.g. adeno and adeno-associated virus vectors, retroviral vectors, Samonella typhi vectors,

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invention, e.g. adeno and adeno-associated virus vectors, retroviral vectors, Salmonella typhi vectors, detoxified anthrax toxin vectors, and the like, will be apparent to those skilled in the art from the description herein.

Furthermore, vaccines in accordance with the invention encompass compositions of one or more of the claimed peptides. A peptide can be present in a vaccine individually. Alternatively, the peptide can exist as a homopolymer comprising multiple copies of the same peptide, or as a heteropolymer of various peptides. Polymers have the advantage of increased immunological reaction and, where different peptide epitopes are used to make up the polymer, the additional ability to induce antibodies and/or CTLs that react with different antigenic determinants of the pathogenic organism or tumor-related peptide targeted for an immune response. The composition can be a naturally occurring region of an antigen or can be prepared, e.g., recombinantly or by chemical synthesis.

Carriers that can be used with vaccines of the invention are well known in the art, and include, e.g., thyroglobulin, albumins such as human serum albumin, tetanus toxoid, polyamino acids such as poly L-lysine, poly L-glutamic acid, influenza, hepatitis B virus core protein, and the like. The vaccines can contain a physiologically tolerable (i.e., acceptable) diluent such as water, or saline, preferably phosphate buffered saline. The vaccines also typically include an adjuvant. Adjuvants such as incomplete Freund's adjuvant, aluminum phosphate, aluminum hydroxide, or alum are examples of materials well known in the art. Additionally, as disclosed herein, CTL responses can be primed by conjugating peptides of the invention to lipids, such as tripalmitory-l-s-glyceryl-systeinlyseryl- serine (P₃CSS).

Upon immunization with a peptide composition in accordance with the invention, via injection, aerosol, oral, transdermal, transmucosal, intrapleural, intrathecal, or other suitable routes, the immune system of the host responds to the vaccine by producing large amounts of CTLs and/or HTLs specific for the desired antigen. Consequently, the host becomes at least partially immune to later infection, or at least partially resistant to developing an ongoing chronic infection, or derives at least some therapeutic benefit when the antigen was tumor-associated.

In some embodiments, it may be desirable to combine the class I peptide components with components that induce or facilitate neutralizing antibody and or helper T cell responses to the target antigen of interest. A preferred embodiment of such a composition comprises class I and class II epitopes in accordance with the invention. An alternative embodiment of such a composition comprises a class I and/or class II epitope in accordance with the invention, along with a cross reactive HTL epitope such as PADRE™ (Epimmune, San Diego, CA) molecule(described e.g., in U.S. Patent Number 5,736,142).

A vaccine of the invention can also include antigen-presenting cells (APC), such as dendritic cells (DC), as a vehicle to present peptides of the invention. Vaccine compositions can be created in vitro, following dendritic cell mobilization and harvesting, whereby loading of dendritic cells occurs in vitro. For example, dendritic cells are transfected, e.g., with a minigene in accordance with the invention, or are pulsed with peptides. The dendritic cell can then be administered to a patient to elicit immune responses in vivo.

Vaccine compositions, either DNA- or peptide-based, can also be administered *in vivo* in combination with dendritic cell mobilization whereby loading of dendritic cells occurs *in vivo*.

Antigenic peptides are used to elicit a CTL and/or HTL response ex vivo, as well. The

10 resulting CTL or HTL cells, can be used to treat chronic infections, or numors in patients that do not respond to other conventional forms of therapy, or will not respond to a therapeutic vaccine peptide or nucleic acid in accordance with the invention. Ex vivo CTL or HTL responses to a particular antigen (infectious or tumor-associated antigen) are induced by incubating in tissue culture the patient's, or genetically compatible, CTL or HTL precursor cells together with a source of antigen-presenting cells (APC), such as dendritic cells, and the appropriate immunogenic peptide. After an appropriate incubation time (typically about 7-28 days), in which the precursor cells are activated and expanded into effector cells,

the cells are infused back into the patient, where they will destroy (CTL) or facilitate destruction (HTL) of their specific target cell (an infected cell or a tumor cell). Transfected dendritic cells may also be used as antigen presenting cells.

The vaccine compositions of the invention may also be used in combination with other

The vaccine compositions of the invention may also be used in commination with other procedures to remove warts or treat HPV infections. Such procedures include cryosurgery, application of caustic agents, electrodessication, surgical excision and laser ablation (Fauci et al. HARRISON'S PRINCIPLES OF INTERNAL MEDICINE, 14th ED., McGraw-Hill Co., Inc, 1998), as well as treatment with antiviral drugs such as interferon-a (see, e.g., Stellato, G., et al., Clin. Diagn. Virol. 7(3):167-72 (1997)) or interferon-inducing drugs such as imiquimod. Topical antimetabolites such a 5-fluorouracil may also be applied.

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In patients with HPV-associated cancer, the vaccine compositions of the invention can also be used in conjunction with other treatments used for cancer, e.g., surgery, chemotherapy, drug therapies, radiation therapies, etc. including use in combination with immune adjuvants such as IL-2, IL-12, GM-CSF, and the like.

Preferably, the following principles are utilized when selecting an array of epitopes for inclusion in a polyepitopic composition for use in a vaccine, or for selecting discrete epitopes to be included in a vaccine and/or to be encoded by nucleic acids such as a minigene. It is preferred that each of the following principles are balanced in order to make the selection. The multiple epitopes to be incorporated in a given vaccine composition may be, but need not be, contiguous in sequence in the native antigen from which the epitopes are derived.

1.) Epitopes are selected which, upon administration, mimic immune responses that have been observed to be correlated with clearance of HPV infection or tumor clearance. For HLA Class I this includes 3-4 epitopes that come from at least one TAA. For HLA Class II a similar rationale is employed; again 3-4 epitopes are selected from at least one TAA (see, e.g., Rosenberg et al., Science

278:1447-1450). Epitopes from one TAA may be used in combination with epitopes from one or more additional TAAs to produce a vaccine that targets tumors with varying expression patterns of frequently-expressed TAAs as described, e.g., in Example 15.

- Epitopes are selected that have the requisite binding affinity established to be
 correlated with immunogenicity: for HLA Class I an IC₃₀ of 500 nM or less, often 200 nM or less; and for Class II an IC₄₀ of 1000 nM or less.
 - 3.) Sufficient supermotif bearing-peptides, or a sufficient array of allele-specific motif-bearing peptides, are selected to give broad population coverage. For example, it is preferable to have at least 80% population coverage. A Monte Carlo analysis, a statistical evaluation known in the art, can be employed to assess the breadth, or redundancy of, population coverage.

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- 4.) When selecting epitopes from cancer-related antigens it is often useful to select analogs because the patient may have developed tolerance to the native epitope. When selecting epitopes for infectious disease-related antigens it is preferable to select either native or analoged epitopes.
- 5.) Of particular relevance are epitopes referred to as "nested epitopes." Nested
 epitopes occur where at least two epitopes overlap in a given peptide sequence. A nested peptide sequence
 can comprise both HLA class I and HLA class II epitopes. When providing nested epitopes, a general
 objective is to provide the greatest number of epitopes per sequence. Thus, an aspect is to avoid providing
 a peptide that is any longer than the amino terminus of the amino terminal epitope and the carboxyl
 terminus of the carboxyl terminal epitope in the peptide. When providing a multi-epitopic sequence, such
 as a sequence comprising nested epitopes, it is generally important to screen the sequence in order to insure
 that it does not have pathological or other deleterious biological properties.
- 6.) If a polyepitopic protein is created, or when creating a minigene, an objective is to generate the smallest peptide that encompasses the epitopes of interest. This principle is similar, if not the same as that employed when selecting a peptide comprising nested epitopes. However, with an artificial polyepitopic peptide, the size minimization objective is balanced against the need to integrate any spacer sequences between epitopes in the polyepitopic protein. Spacer amino acid residues can, for example, be introduced to avoid junctional epitopes (an epitope recognized by the immune system, not present in the target antigen, and only created by the man-made juxtaposition of epitopes), or to facilitate cleavage between epitopes and thereby enhance epitope presentation. Junctional epitopes are generally to be avoided because the recipient may generate an immune response to that non-native epitope. Of particular concern is a junctional epitope that is a "dominant epitope." A dominant epitope may lead to such a zealous response that immune responses to other epitopes are diminished or suppressed.
 - 7.) In cases where the sequences of multiple variants of the same target protein are available, potential peptide epitopes can also be selected on the basis of their conservancy. For example, a criterion for conservancy may define that the entire sequence of an HLA class I binding peptide or the entire 9-mer core of a class II binding peptide be conserved in a designated percentage of the sequences evaluated for a specific protein antigen.
- 8.) When selecting an array of epitopes of an infectious agent, it is preferred that at least some of the epitopes are derived from early and late proteins. The early proteins of HPV are expressed when the virus is replicating, either following acute or dormant infection. Therefore, it is

particularly preferred to use epitopes from early stage proteins to alleviate disease manifestations at the earliest stage possible.

III.K.1. Minigene Vaccines

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5 A number of different approaches are available which allow simultaneous delivery of multiple epitopes. Nucleic acids encoding the peptides of the invention are a particularly useful embodiment of the invention. Epitopes for inclusion in a minigene are preferably selected according to the guidelines set forth in the previous section. A preferred means of administering nucleic acids encoding the peptides of the invention uses minigene constructs encoding a peptide comprising one or multiple epitopes of the invention.

The use of multi-epitope minigenes is described below and in, e.g., co-pending application U.S.S.N. 09311,784; Ishioka et al., J. Immunol. 162:3915-3925, 1999; An, L. and Whitton, J. L., J. Virol. 71:2292, 1997; Thomson, S. A. et al., J. Immunol. 157:822, 1996; Whitton, J. L. et al., J. Virol. 67:348, 1993; Hanke, R. et al., Vaccine 16:426, 1998. For example, a multi-epitope DNA plasmid

15 encoding supermotif- and/or motif-bearing epitopes derived from multiple regions of one or more HPV antigens, the PADRE universal helper T cell epitope (or multiple HTL epitopes from HPV antigens), and an endoplasmic reticulum-translocating signal sequence can be engineered. A vaccine may also comprise epitopes that are derived from other TAAs.

The immunogenicity of a multi-epitopic minisene can be tested in transgenic mice to evaluate the magnitude of CTL induction responses against the epitopes tested. Further, the immunogenicity of DNA-encoded epitopes in vivo can be correlated with the in vitro responses of specific CTL lines against target cells transfected with the DNA plasmid. Thus, these experiments can show that the minisgene serves to both: 1.) generate a CTL response and 2.) that the induced CTLs recognized cells expressing the encoded epitopes.

For example, to create a DNA sequence encoding the selected epitopes (minigene) for expression in human cells, the amino acid sequences of the epitopes may be reverse translated. A human codon usage table can be used to guide the codon choice for each amino acid. These epitope-encoding DNA sequences may be directly adjoined, so that when translated, a continuous polypeptide sequence is created. To optimize expression and/or immunogenicity, additional elements can be incorporated into the minigene design. Examples of amino acid sequences that can be reverse translated and included in the minigene sequence include: HLA class I epitopes, HLA class II epitopes, a ubiquitination signal sequence, and/or an endoplasmic reticulum targeting signal. In addition, HLA presentation of CTL and HTL epitopes may be improved by including synthetic (e.g. poly-alanine) or naturally-occurring flanking sequences adjacent to the CTL or HTL epitopes; these larger peptides comprising the epitope(s) are within the scope of the invention.

The minigene sequence may be converted to DNA by assembling oligonucleotides that encode the plus and minus strands of the minigene. Overlapping oligonucleotides (30-100 bases long) may be synthesized, phosphorylated, purified and annealed under appropriate conditions using well known techniques. The ends of the oligonucleotides can be joined, for example, using T4 DNA ligase. This synthetic minigene, encoding the epitope polypeptide, can then be cloned into a desired expression vector.

Standard regulatory sequences well known to those of skill in the art are preferably included in the vector to ensure expression in the target cells. Several vector elements are desirable: a promoter with a down-stream cloning site for minigene insertion, a polyadenylation signal for efficient transcription termination; an E. coli origin of replication; and an E. coli selectable market (e.g. ampicillin or kanamycin resistance). Numerous promoters can be used for this purpose, e.g., the human cytomegalovirus (hCMV) promoter. See, e.g., U.S. Patent Nos. 5, 80, 859 and 5, 589, 466 for other suitable promoter sequences.

Additional vector modifications may be desired to optimize minigene expression and immunogenicity. In some cases, introns are required for efficient gene expression, and one or more synthetic or naturally-occurring introns could be incorporated into the transcribed region of the minigene. The inclusion of mRNA stabilization sequences and sequences for replication in mammalian cells may also be considered for increasing minigene expression.

Once an expression vector is selected, the minigene is cloned into the polylinker region downstream of the promoter. This plasmid is transformed into an appropriate E. coli strain, and DNA is prepared using standard techniques. The orientation and DNA sequence of the minigene, as well as all other elements included in the vector, are confirmed using restriction mapping and DNA sequence analysis. Bacterial cells harboring the correct plasmid can be stored as a master cell bank and a working cell bank.

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In addition, immunostimulatory sequences (ISSs or CpGs) appear to play a role in the immunogenicity of DNA vaccines. These sequences may be included in the vector, outside the minigene coding sequence, if desired to enhance immunogenicity.

In some embodiments, a bi-cistronic expression vector which allows production of both
the minigene-encoded epitopes and a second protein (included to enhance or decrease immunogenicity) can
be used. Examples of proteins or polypeptides that could beneficially enhance the immune response if coexpressed include cytokines (e.g., IL-2, IL-12, GM-CSF), cytokine-inducing molecules (e.g., LeIF),
25 costimulatory molecules, or for HTL responses, pan-DR binding proteins (PADRE™, Epirmmune, San
Diego, CA). Helper (HTL) epitopes can be joined to intracellular targeting signals and expressed separately
from expressed CTL epitopes; this allows direction of the HTL epitopes to a cell compartment different
than that of the CTL epitopes; If required, this could facilitate more efficient entry of HTL epitopes into the
HLA class II pathway, thereby improving HTL induction. In contrast to HTL or CTL induction,
30 specifically decreasing the immune response by co-expression of immunosuppressive molecules (e.g. TGF8) may be beneficial in certain diseases.

Therapeutic quantities of plasmid DNA can be produced for example, by fermentation in E. coli, followed by purification. Aliquots from the working cell bank are used to inoculate growth medium, and grown to saturation in shaker flasks or a bioreactor according to well known techniques. Plasmid DNA can be purified using standard bioseparation technologies such as solid phase amionexchange resins supplied by QIAGEN, Inc. (Valencia, California). If required, supercoiled DNA can be isolated from the open circular and linear forms using gel electrophoresis or other methods.

Purified plasmid DNA can be prepared for injection using a variety of formulations. The simplest of these is reconstitution of lyophilized DNA in sterile phosphate-buffer saline (PBS). This approach, known as "naked DNA," is currently being used for intramuscular (IM) administration in clinical

trials. To maximize the immunotherapeutic effects of minigene DNA vaccines, an alternative method for formulating purified plasmid DNA may be desirable. A variety of methods have been described, and new techniques may become available. Cationic lipids, glycolipids, and fusogenic liposomes can also be used in the formulation (see, e.g., as described by WO 93/24640; Mannino & Gould-Fogerite, BioTechniques 6(7): 682 (1988); U.S. Pat No. 5,279,833; WO 91/06309; and Felgner, et al., Proc. Nat'l Acad. Sci. USA 84:7413 (1987). In addition, peptides and compounds referred to collectively as protective, interactive, non-condensing compounds (PINC) could also be complexed to purified plasmid DNA to influence variables such as stability, intramuscular dispersion, or trafficking to specific organs or cell types.

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Target cell sensitization can be used as a functional assay for expression and HLA class I

or presentation of minigene-encoded CTL epitopes. For example, the plasmid DNA is introduced into a
mammalian cell line that is suitable as a target for standard CTL chromium release assays. The transfection
method used will be dependent on the final formulation. Electroporation can be used for "naked" DNA,
whereas cationic lipids allow direct in vitro transfection. A plasmid expressing green fluorescent protein
(GFP) can be co-transfected to allow enrichment of transfected cells using fluorescence activated cell
sorting (FACS). These cells are then chromium-51 ("Or) labeled and used as target cells for epitopespecific CTL lines; cytolysis, detected by "1cr release, indicates both production of, and HLA presentation
of, minigene-encoded CTL epitopes. Expression of HTL epitopes may be evaluated in an analogous

In vivo immunogenicity is a second approach for functional testing of minigene DNA

formulations. Transgenic mice expressing appropriate human HLA proteins are immunized with the DNA
product. The dose and route of administration are formulation dependent (e.g., IM for DNA in PBS,
intraperitoneal (i.p.) for lipid-complexed DNA). Twenty-one days after immunization, splenocytes are
harvested and restimulated for one week in the presence of peptides encoding each epitope being tested.
Thereafter, for CTL effector cells, assays are conducted for cytolysis of peptide-loaded, ³¹Cr-labeled target
cells using standard techniques. Lysis of target cells that were sensitized by HLA loaded with peptide
epitopes, corresponding to minigene-encoded epitopes, demonstrates DNA vaccine function for in vivo
induction of CTLs. Immunogenicity of HTL epitopes is evaluated in transgenic mice in an analogous
manner.

Alternatively, the nucleic acids can be administered using ballistic delivery as described, for instance, in U.S. Patent No. 5,204,253. Using this technique, particles comprised solely of DNA are administered. In a further alternative embodiment, DNA can be adhered to particles, such as gold particles. Minigenes can also be delivered using other bacterial or viral delivery systems well known in the art, e.g., an expression construct encoding epitopes of the invention can be incorporated into a viral vector such as vaccinia.

III.K.2. Combinations of CTL Peptides with Helper Peptides

manner using assays to assess HTL activity.

Vaccine compositions comprising CTL peptides of the invention can be modified to provide desired attributes, such as improved serum half life, broadened population coverage or enhanced immunogenicity.

For instance, the ability of a peptide to induce CTL activity can be enhanced by linking the peptide to a sequence which contains at least one epitope that is capable of inducing a T helper cell response. The use of T helper epitopes in conjunction with CTL epitopes to enhance immunogenicity is illustrated, for example, in the co-pending applications U.S.S.N. 08/820,360, U.S.S.N. 08/197,484, and U.S.S.N. 08/8444 234.

Although a CTL peptide can be directly linked to a T helper peptide, often CTL epitope/HTL epitope conjugates are linked by a spacer molecule. The spacer is typically comprised of relatively small, neutral molecules, such as amino acids or amino acid mimetics, which are substantially uncharged under physiological conditions. The spacers are typically selected from, e.g., Ala, Gly, or other neutral spacers of nonpolar amino acids or neutral polar amino acids. It will be understood that the optionally present spacer need not be comprised of the same residues and thus may be a hetero- or homoligomer. When present, the spacer will usually be at least one or two residues, more usually three to six residues and sometimes 10 or more residues. The CTL peptide epitope can be linked to the T helper peptide epitope either directly or via a spacer either at the amino or carboxy terminus of the CTL peptide. The amino terminus of either the immunogenic peptide or the T helper peptide may be acylated.

In certain embodiments, the T helper peptide is one that is recognized by T helper cells present in the majority of the population. This can be accomplished by selecting peptides that bind to many, most, or all of the HLA class II molecules. These are known as "loosely HLA-restricted" or "promiscuous" T helper sequences. Examples of amino acid sequences that are promiscuous include sequences from antigens such as tetanus toxoid at positions 830-843 (QYIKANSKFIGITE; SEQ ID NO: 51484), Plasmodium falciparum circumsporozoite (CS) protein at positions 378-398 (DIEKKIAKMEKASSVFNVVNS; SEQ ID NO: 51485), and Streptococcus 18kD protein at positions 116 (GAVDSILGGVATYGAA; SEQ ID NO: 51486). Other examples include peptides bearing a DR 1-4-7 supermotif, or either of the DR3 motifs.

Alternatively, it is possible to prepare synthetic peptides capable of stimulating Theiper lymphocytes, in a loosely HLA-restricted fashion, using amino acid sequences not found in nature (see, e.g., PCT publication WO 95/07707). These synthetic compounds called Pan-DR-binding epitopes (e.g., PADRE™, Epimmune, Inc., San Diego, CA) are designed to most preferrably bind most HLA-DR (human HLA class II) molecules. For instance, a pan-DR-binding epitope peptide having the formula: aKXVAAWTLKAAa, where "X" is either cyclohexylalanine, phenylalanine, or tyrosine, and a is either Dalanine or L-alanine, has been found to bind to most HLA-DR alleles, and to stimulate the response of T helper lymphocytes from most individuals, regardless of their HLA type. An alternative of a pan-DR binding epitope comprises all "L" natural amino acids and can be provided in the form of nucleic acids that encode the epitope.

HTL peptide epitopes can also be modified to alter their biological properties. For example, they can be modified to include p-amino acids to increase their resistance to proteases and thus extend their serum half life, or they can be conjugated to other molecules such as lipids, proteins, carbohydrates, and the like to increase their biological activity. For example, a Thelper peptide can be conjugated to one or more palmitic acid chains at either the amino or carboxyl termini.

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III.K.3. Combinations of CTL Peptides with T Cell Priming Agents

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In some embodiments it may be desirable to include in the pharmaceutical compositions of the invention at least one component which primes cytotoxic T lymphocytes. Lipids have been identified as agents capable of priming CTL in vivo against viral antigens. For example, palmitic acid residues can be attached to the ϵ -and α - amino groups of a lysine residue and then linked, ϵ , ϵ , via one or more linking residues such as Gly, Gly-Gly-, Ser, Ser-Ser, or the like, to an immunogenic peptide. The lipidated peptide can then be administered either directly in a micelle or particle, incorporated into a liposome, or emulsified in an adjuvant, ϵ , ϵ , incomplete Freund's adjuvant. In a preferred embodiment, a particularly effective immunogenic composition comprises palmitic acid attached to ϵ - and α - amino groups of Lys, which is attached via linkage, ϵ , ϵ , Ser-Ser, to the amino terminus of the immunogenic peptide.

As another example of lipid priming of CTL responses, E. coli lipoproteins, such as tripalmitoyl-S-glycerylcysteinlyseryl-serine (P₂CSS) can be used to prime virus specific CTL when covalently attached to an appropriate peptide (see, e.g., Deres, et al., Nature 342:561, 1989). Peptides of the invention can be coupled to P₂CSS, for example, and the lipopeptide administered to an individual to specifically prime a CTL response to the target antigen. Moreover, because the induction of neutralizing antibodies can also be primed with P₂CSS-conjugated epitopes, two such compositions can be combined to more effectively elicit both humoral and cell-mediated responses.

CTL and/or HTL peptides can also be modified by the addition of amino acids to the termini of a peptide to provide for ease of linking peptides one to another, for coupling to a carrier support or larger peptide, for modifying the physical or chemical properties of the peptide or oligopeptide, or the like. Amino acids such as tyrosine, cysteine, lysine, glutamic or aspartic acid, or the like, can be introduced at the C- or N-terminus of the peptide or oligopeptide, particularly class I peptides. However, it is to be noted that modification at the carboxyl terminus of a CTL epitope may, in some cases, alter binding characteristics of the peptide. In addition, the peptide or oligopeptide sequences can differ from the natural sequence by being modified by terminal-NH₃ acylation, e.g., by alkanoyl (CI-C20) or thioglycolyl acetylation, terminal-carboxyl amidation, e.g., ammonia, methylamine, etc. In some instances these modifications may provide sites for linking to a support or other molecule.

IV.J.4. Vaccine Compositions Comprising DC Pulsed with CTL and/or HTL Peptides

An embodiment of a vaccine composition in accordance with the invention comprises ex vivo administration of a cocktail of epitope-bearing peptides to PBMC, or isolated DC therefrom, from the patient's blood. A pharmaceutical to facilitate harvesting of DC can be used, such as Progenipoietin (Monsanto, St. Louis, MO) or GM-CSF/IL-4. After pulsing the DC with peptides and prior to reinfusion into patients, the DC are washed to remove unbound peptides. In this embodiment, a vaccine comprises peptide-pulsed DCs which present the pulsed peptide epitopes complexed with HLA molecules on their surfaces.

The DC can be pulsed ex vivo with a cocktail of peptides, some of which stimulate CTL responses to one or more HPV antigens of interest. Optionally, a helper T cell (HTL) peptide such as a PADRE family molecule, can be included to facilitate the CTL response. Thus, a vaccine in accordance

with the invention, preferably comprising epitopes from multiple HPV antigens, is used to treat HPV infection or cancer resulting from HPV infection.

III.L. Administration of Vaccines for Therapeutic or Prophylactic Purposes

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5 The peptides of the present invention and pharmaceutical and vaccine compositions of the invention are typically used to treat and/or prevent cancer associated with HPV infection. Vaccine compositions containing the peptides of the invention are administered to a patient infected with HPV or to an individual susceptible to, or otherwise at tisk for, HPV infection to elicit an immune response against HPV antigens and thus enhance the patient's own immune response capabilities.

As noted above, peptides comprising CTL and/or HTL epitopes of the invention induce immune responses when presented by HLA molecules and contacted with a CTL or HTL specific for an epitope comprised by the peptide. The peptides (or DNA encoding them) can be administered individually or as fusions of one or more peptide sequences. The manner in which the peptide is contacted with the CTL or HTL is not critical to the invention. For instance, the peptide can be contacted with the CTL or HTL either in vivo or in vitro. If the contacting occurs in vivo, the peptide itself can be administered to the patient, or other vehicles, e.g., DNA vectors encoding one or more peptides, viral vectors encoding the peptide(s), liposomes and the like, can be used, as described herein.

When the peptide is contacted in vitro, the vaccinating agent can comprise a population of cells, e.g., peptide-pulsed dendritic cells, or HPV-specific CTLs, which have been induced by pulsing antigen-presenting cells in vitro with the peptide or by transfecting antigen-presenting cells with a minigene of the invention. Such a cell population is subsequently administered to a patient in a therapeutically effective dose.

In therapeutic applications, peptide and/or nucleic acid compositions are administered to a patient in an amount sufficient to elicit an effective CTL and/or HTL response to the virus antigen and to cure or at least partially arrest or slow symptoms and/or complications. An amount adequate to accomplish this is defined as "therapeutically effective dose." Amounts effective for this use will depend on, e.g., the particular composition administered, the manner of administration, the stage and severity of the disease being treated, the weight and general state of health of the patient, and the judgment of the prescribing physician.

For pharmaceutical compositions, the immunogenic peptides of the invention, or DNA encoding them, are generally administered to an individual already infected with HPV. The peptides or DNA encoding them can be administered individually or as fusions of one or more peptide sequences. HPV-infected patients, with or without neoplasia, can be treated with the immunogenic peptides separately or in conjunction with other treatments, such as surgery, as appropriate.

For therapeutic use, administration should generally begin at the first diagnosis of HPV infection or HPV-associated cancer. This is followed by boosting doses until at least symptoms are substantially abated and for a period thereafter. The embodiment of the vaccine composition (i.e., including, but not limited to embodiments such as peptide cocktails, polyepitopic polypeptides, minigenes, or TAA-specific CTLs or pulsed dendritic cells) delivered to the patient may vary according to the stage of the disease or the patient's health status. For example, in a patient with a tumor that expresses HPV

antigens, a vaccine comprising HPV-specific CTL may be more efficacious in killing tumor cells in patient with advanced disease than alternative embodiments.

Where susceptible individuals are identified prior to or during infection, the composition can be targeted to them, thus minimizing the need for administration to a larger population. Susceptible populations include those individuals who are sexually active.

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The peptide or other compositions used for the treatment or prophylaxis of HPV infection can be used, e.g., in persons who have not manifested symptoms, e.g., genital warts or neoplastic growth. In this context, it is generally important to provide an amount of the peptide epitope delivered by a mode of administration sufficient to effectively stimulate a cytotoxic T cell response; compositions which stimulate helper T cell responses can also be given in accordance with this embodiment of the invention.

The dosage for an initial therapeutic immunization generally occurs in a unit dosage range where the lower value is about 1, 5, 50, 500, or 1,000 µg and the higher value is about 10,000; 20,000; 30,000; or 50,000 µg. Dosage values for a human typically range from about 500 µg to about 50,000 µg per 70 kilogram patient. Boosting dosages of between about 10 µg to about 50,000 µg of peptide pursuant to a boosting regimen over weeks to months may be administered depending upon the patient's response and condition as determined by measuring the specific activity of CTL and HTL obtained from the patient's blood. Administration should continue until at least clinical symptoms or laboratory tests indicate that the viral infection, or neoplasia, has been eliminated or reduced and for a period thereafter. The dosages, routes

In certain embodiments, the peptides and compositions of the present invention are employed in serious disease states, that is, life-threatening or potentially life threatening situations. In such cases, as a result of the minimal amounts of extraneous substances and the relative nontoxic nature of the peptides in preferred compositions of the invention, it is possible and may be felt desirable by the treating physician to administer substantial excesses of these peptide compositions relative to these stated dosage amounts.

of administration, and dose schedules are adjusted in accordance with methodologies known in the art.

The vaccine compositions of the invention can also be used purely as prophylactic agents.

Generally the dosage for an initial prophylactic immunization generally occurs in a unit dosage range where the lower value is about 1, 5, 50, 500, or 1000 µg and the higher value is about 10,000; 20,000; 30,000; or 50,000 µg. Dosage values for a human typically range from about 500 µg to about 50,000 µg per 70 kilogram patient. This is followed by boosting dosages of between about 1.0 µg to about 50,000 µg of peptide administered at defined intervals from about four weeks to six months after the initial administration of vaccine. The immunogenicity of the vaccine can be assessed by measuring the specific activity of CTL and HTL obtained from a sample of the patient's blood.

The pharmaceutical compositions for therapeutic treatment are intended for parenteral, topical, oral, intrathecal, or local (e.g. as a cream or topical ointment) administration. Preferably, the pharmaceutical compositions are administered parentally, e.g., intravenously, subcutaneously, intradermally, or intramuscularly. Thus, the invention provides compositions for parenteral administration which comprise a solution of the immunogenic peptides dissolved or suspended in an acceptable carrier, preferably an aqueous carrier. A variety of aqueous carriers may be used, e.g., water, buffered water, 0.8% saline, 0.3% glycine, hyaluronic acid and the like. These compositions may be sterilized by conventional,

well known sterilization techniques, or may be sterile filtered. The resulting aqueous solutions may be packaged for use as is, or lyophilized, the lyophilized preparation being combined with a sterile solution prior to administration. The compositions may contain pharmaceutically acceptable auxiliary substances as required to approximate physiological conditions, such as pH-adjusting and buffering agents, tonicity adjusting agents, wetting agents, preservatives, and the like, for example, sodium acetate, sodium lactate, sodium chloride, potassium chloride, calcium chloride, sorbian monolaurate, triethanolamine oleate, etc.

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The concentration of peptides of the invention in the pharmaceutical formulations can vary widely, i.e., from less than about 0.1%, usually at or at least about 2% to as much as 20% to 50% or more by weight, and will be selected primarily by fluid volumes, viscosities, etc., in accordance with the particular mode of administration selected.

A human unit dose form of the peptide composition is typically included in a pharmaceutical composition that comprises a human unit dose of an acceptable carrier, preferably an aqueous carrier, and is administered in a volume of fluid that is known by those of skill in the art to be used for administration of such compositions to humans (see, e.g., Remington's Pharmaceutical Sciences, 17th Edition, A. Gennaro, Editor, Mack Publising Co., Easton, Pennsylvania, 1985).

The peptides of the invention, and/or nucleic acids encoding the peptides, can also be administered via liposomes, which may also serve to target the peptides to a particular tissue, such as lymphoid tissue, or to target selectively to infected cells, as well as to increase the half-life of the peptide composition. Liposomes include emulsions, foams, micelles, insoluble monolayers, liquid crystals, phospholipid dispersions, lamellar layers and the like. In these preparations, the peptide to be delivered is incorporated as part of a liposome, alone or in conjunction with a molecule which binds to a receptor prevalent among lymphoid cells, such as monoclonal antibodies which bind to the CD45 antigen, or with other therapeutic or immunogenic compositions. Thus, liposomes either filled or decorated with a desired peptide of the invention can be directed to the site of lymphoid cells, where the liposomes then deliver the peptide compositions. Liposomes for use in accordance with the invention are formed from standard vesicle-forming lipids, which generally include neutral and negatively charged phospholipids and a sterol, such as cholesterol. The selection of lipids is generally guided by consideration of, e.g., liposome size, acid lability and stability of the liposomes in the blood stream. A variety of methods are available for preparing liposomes, as described in, e.g., Szoka, et al., Ann. Rev. Biophys. Bioeng. 9.467 (1980), and U.S. Patent Nos. 4.253,871, 4,501,728, 4,337,028, and 5,019,369.

For targeting cells of the immune system, a ligand to be incorporated into the liposome can include, e.g., antibodies or fragments thereof specific for cell surface determinants of the desired immune system cells. A liposome suspension containing a peptide may be administered intravenously, locally, topically, etc. in a dose which varies according to, inter alia, the manner of administration, the peptide being delivered, and the stage of the disease being treated.

For solid compositions, conventional nontoxic solid carriers may be used which include, for example, pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharin, talcum, cellulose, glucose, sucrose, magnesium carbonate, and the like. For oral administration, a pharmaceutically acceptable nontoxic composition is formed by incorporating any of the normally

employed excipients, such as those carriers previously listed, and generally 10-95% of active ingredient, that is, one or more peptides of the invention, and more preferably at a concentration of 25%-75%.

For aerosol administration, the immunogenic peptides are preferably supplied in finely divided form along with a surfactant and propellant. Typical percentages of peptides are 0.01%-20% by weight, preferably 1%-10%. The surfactant must, of course, be nontoxic, and preferably soluble in the propellant. Representative of such agents are the esters or partial esters of farty acids containing from 6 to 22 carbon atoms, such as caproic, octanoic, lauric, palmitic, stearic, linoleic, linolenic, olesteric and oleic acids with an aliphatic polyhydric alcohol or its cyclic anhydride. Mixed esters, such as mixed or natural glycerides may be employed. The surfactant may constitute 0.1%-20% by weight of the composition, preferably 0.25-5%. The balance of the composition is ordinarily propellant. A carrier can also be included, as desired, as with, e.g., lecithin for intranasal delivery.

III.M. HLA EXPRESSION: IMPLICATIONS FOR T CELL-BASED IMMUNOTHERAPY Disease progression in cancer and infectious disease

It is well recognized that a dynamic interaction between exists between host and disease, both in the cancer and infectious disease settings. In the infectious disease setting, it is well established that pathogens evolve during disease. The strains that predominate early in HIV infection are different from the ones that are associated with AIDS and later disease stages (NS versus S strains). It has long been hypothesized that pathogen forms that are effective in establishing infection may differ from the ones most effective in terms of replication and chronicity.

Similarly, it is widely recognized that the pathological process by which an individual succumbs to a neoplastic disease is complex. During the course of disease, many changes occur in cancer cells. The timor accumulates alterations which are in part related to dysfunctional regulation of growth and differentiation, but also related to maximizing its growth potential, escape from drug treatment and/or the body's immunosurveillance. Neoplastic disease results in the accumulation of several different biochemical alterations of cancer cells, as a function of disease progression. It also results in significant levels of intra- and inter- cancer heterogeneity, particularly in the late, metastatic stage.

Familiar examples of cellular alterations affecting treatment outcomes include the outgrowth of radiation or chemotherapy resistant tumors during the course of therapy. These examples parallel the emergence of drug resistant viral strains as a result of aggressive chemotherapy, e.g., of chronic HBV and HIV infection, and the current resurgence of drug resistant organisms that cause Tuberculosis and Malaria. It appears that significant heterogeneity of responses is also associated with other approaches to cancer therapy, including anti-angiogenesis drugs, passive antibody immunotherapy, and active T cell-based immunotherapy. Thus, in view of such phenomena, epitopes from multiple disease-related antigens can be used in vaccines and therapeutics thereby counteracting the ability of diseased cells to mutate and escape treatment.

The interplay between disease and the immune system

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One of the main factors contributing to the dynamic interplay between host and disease is

40 the immune response mounted against the pathogen, infected cell, or malignant cell. In many conditions

such immune responses control the disease. Several animal model systems and prospective studies of natural infection in humans suggest that immune responses against a pathogen can control the pathogen, prevent progression to severe disease and/or eliminate the pathogen. A common theme is the requirement for a multispecific T cell response, and that narrowly focused responses appear to be less effective. These observations guide skilled artisan as to embodiments of methods and compositions of the present invention that provide for a broad immune response.

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In the cancer setting there are several findings that indicate that immune responses can impact neoplastic growth:

First, the demonstration in many different animal models, that anti-tumor T cells, restricted by MHC class I, can prevent or treat tumors.

Second, encouraging results have come from immunotherapy trials.

Third, observations made in the course of natural disease correlated the type and composition of T cell infiltrate within tumors with positive clinical outcomes (Coulie PG, et al. Antitumor immunity at work in a melanoma patient In <u>Advances in Cancer Research</u>, 213-242, 1999).

- Finally, tumors commonly have the ability to mutate, thereby changing their 15 immunological recognition. For example, the presence of monospecific CTL was also correlated with control of tumor growth, until antigen loss emerged (Riker A, et al., Immune selection after antigen-specific immunotherapy of melanoma Surgery, Aug: 126(2):112-20, 1999; Marchand M, et al., Tumor regressions observed in patients with metastatic melanoma treated with an antigenic peptide encoded by gene MAGE-3 and presented by HLA-A1 Int. J. Cancer 80(2):219-30, Jan. 18, 1999). Similarly, loss of beta 2 20 microglobulin was detected in 5/13 lines established from melanoma patients after receiving immunotherapy at the NCI (Restifo NP, et al., Loss of functional Beta2 - microglobulin in metastatic melanomas from five patients receiving immunotherapy Journal of the National Cancer Institute, Vol. 88 (2), 100-108, Jan. 1996). It has long been recognized that HLA class I is frequently altered in various tumor types. This has led to a hypothesis that this phenomenon might reflect immune pressure exerted on 25 the tumor by means of class I restricted CTL. The extent and degree of alteration in HLA class I expression appears to be reflective of past immune pressures, and may also have prognostic value (van Duinen SG, et al., Level of HLA antigens in locoregional metastases and clinical course of the disease in patients with melanoma Cancer Research 48, 1019-1025, Feb. 1988; Möller P, et al., Influence of major histocompatibility complex class I and II antigens on survival in colorectal carcinoma Cancer Research 51, 30 729-736, Jan. 1991). Taken together, these observations provide a rationale for immunotherapy of cancer and infectious disease, and suggest that effective strategies need to account for the complex series of pathological changes associated with disease.
- The three main types of alterations in HLA expression in tumors and their functional significance

 The level and pattern of expression of HLA class I antigens in tumors has been studied in many different tumor types and alterations have been reported in all types of tumors studied. The molecular mechanisms underlining HLA class I alterations have been demonstrated to be quite heterogeneous. They include alterations in the TAP/processing pathways, mutations of β2-microglobulin and specific HLA heavy chains, alterations in the regulatory elements controlling over class I expression and loss of entire

chromosome sections. There are several reviews on this topic, see, e.g., : Garrido F, et al., Natural history of HLA expression during tumour development Immunol Today 14(10):491-499, 1993; Kaklamanis L, et al., Loss of HLA class-I alleles, heavy chains and \(\beta 2\)-microglobulin in colorectal cancer Int. J. Cancer, 51(3):379-85, May 28,1992. There are three main types of HLA Class I alteration (complete loss, allelespecific loss and decreased expression). The functional significance of each alteration is discussed separately:

Complete loss of HLA expression

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Complete loss of HLA expression can result from a variety of different molecular mechanisms, reviewed in (Algarra I, et al., The HLA crossroad in tumor immunology Human Immunology 10 61, 65-73, 2000; Browning M, et al., Mechanisms of loss of HLA class I expression on colorectal tumor cells Tissue Antigens 47:364-371, 1996; Ferrone S, et al., Loss of HLA class I antigens by melanoma cells: molecular mechanisms, functional significance and clinical relevance Immunology Today, 16(10): 487-494, 1995; Garrido F, et al., Natural history of HLA expression during tumour development Immunology Today 14(10):491-499, 1993; Tait, BD, HLA Class I expression on human cancer cells: Implications for effective 15 immunotherapy Hum Immunol 61, 158-165, 2000). In functional terms, this type of alteration has several important implications.

While the complete absence of class I expression will eliminate CTL recognition of those tumor cells, the loss of HLA class I will also render the tumor cells extraordinary sensitive to lysis from NK cells (Ohnmacht, GA, et al., Heterogeneity in expression of human leukocyte antigens and melanomaassociated antigens in advanced melanoma J Cellular Phys 182:332-338, 2000; Liunggren HG, et al., Host resistance directed selectively against H-2 deficient lymphoma variants: Analysis of the mechanism J. Exp. Med., Dec 1;162(6):1745-59, 1985; Maio M, et al., Reduction in susceptibility to natural killer cellmediated lysis of human FO-1 melanoma cells after induction of HLA class I antigen expression by transfection with B2m gene J. Clin. Invest. 88(1):282-9, July 1991; Schrier PI, et al., Relationship between myc oncogene activation and MHC class I expression Adv. Cancer Res., 60:181-246, 1993).

The complementary interplay between loss of HLA expression and gain in NK sensitivity is exemplified by the classic studies of Coulie and coworkers (Coulie, PG, et al., Antitumor immunity at work in a melanoma patient. In Advances in Cancer Research, 213-242, 1999) which described the 30 evolution of a patient's immune response over the course of several years. Because of increased sensitivity to NK lysis, it is predicted that approaches leading to stimulation of innate immunity in general and NK activity in particular would be of special significance. An example of such approach is the induction of large amounts of dendritic cells (DC) by various hematopoietic growth factors, such as Flt3 ligand or ProGP. The rationale for this approach resides in the well known fact that dendritic cells produce large amounts of IL-12, one of the most potent stimulators for innate immunity and NK activity in particular. Alternatively, IL-12 is administered directly, or as nucleic acids that encode it. In this light, it is interesting to note that Flt3 ligand treatment results in transient tumor regression of a class I negative prostate murine cancer model (Ciavarra RP, et al., Flt3-Ligand induces transient tumor regression in an ectopic treatment model of major histocompatibility complex-negative prostate cancer Cancer Res 60:2081-84, 2000). In this context, specific anti-tumor vaccines in accordance with the invention synergize with these types of

hematopoietic growth factors to facilitate both CTL and NK cell responses, thereby appreciably impairing a cell's ability to mutate and thereby escape efficacious treatment. Thus, an embodiment of the present invention comprises a composition of the invention together with a method or composition that augments functional activity or numbers of NK cells. Such an embodiment can comprise a protocol that provides a composition of the invention sequentially with an NK-inducing modality, or contemporaneous with an NK-inducing modality.

Secondly, complete loss of HLA frequently occurs only in a fraction of the tumor cells, while the remainder of tumor cells continue to exhibit normal expression. In functional terms, the tumor would still be subject, in part, to direct attack from a CTL response; the portion of cells lacking HLA subject to an NK response. Even if only a CTL response were used, destruction of the HLA expressing fraction of the tumor has dramatic effects on survival times and quality of life.

It should also be noted that in the case of heterogeneous HLA expression, both normal HLA-expressing as well as defective cells are predicted to be susceptible to immune destruction based on "bystander effects." Such effects were demonstrated, e.g., in the studies of Rosendahl and colleagues that investigated in vivo mechanisms of action of antibody targeted superantigens (Rosendahl A, et al., Perforin and IFN-gamma are involved in the antitumor effects of antibody-targeted superantigens J. Immunol. 160(11):5309-13, June 1, 1998). The bystander effect is understood to be mediated by cytokines elicited from, e.g., CTLs acting on an HLA-bearing target cell, whereby the cytokines are in the environment of other diseased cells that are concomitantly killed.

Allele-specific loss

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One of the most common types of alterations in class I molecules is the selective loss of certain alleles in individuals heterozygous for HLA. Allele-specific alterations might reflect the tumor adaptation to immune pressure, exerted by an immunodominant response restricted by a single HLA restriction element. This type of alteration allows the tumor to retain class I expression and thus escape NK cell recognition, yet still be susceptible to a CTL-based vaccine in accordance with the invention which comprises epitopes corresponding to the remaining HLA type. Thus, a practical solution to overcome the potential hurdle of allele-specific loss relies on the induction of multispecific responses. Just as the inclusion of multiple disease-associated antigens in a vaccine of the invention guards against mutations that yield loss of a specific disease antigens, simultaneously targeting multiple HLA specificities and multiple disease-related antigens prevents disease escape by allele-specific losses.

Decrease in expression (allele-specific or not)

The sensitivity of effector CTL has long been demonstrated (Brower, RC, et al., Minimal requirements for peptide mediated activation of CD8+ CTL Mol. Immunol., 31;1285-93, 1994; Chriustnick, ET, et al. Low numbers of MHC class I-peptide complexes required to trigger a T cell response Nature 352:67-70, 1991; Sykulev, Y, et al., Evidence that a single peptide-MHC complex on a target cell can elicit a cytolytic T cell response Immunity, 4(6):565-71, June 1996). Even a single peptide/MHC complex can result in tumor cells lysis and release of anti-tumor lymphokines. The biological significance of decreased HLA expression and possible tumor escape from immune recognition is not fully known. Nevertheless, it

has been demonstrated that CTL recognition of as few as one MHC/peptide complex is sufficient to lead to tumor cell lysis.

Further, it is commonly observed that expression of HLA can be upregulated by gamma IFN, commonly secreted by effector CTL. Additionally, HLA class I expression can be induced in vivo by both alpha and beta IFN (Halloran, et al. Local T cell responses induce widespread MHC expression. J Immunol 148:3837, 1992; Pestka, S, et al., Interferons and their actions Annu. Rev. Biochem. 56:727-77, 1987). Conversely, decreased levels of HLA class I expression also render cells more susceptible to NK lysis.

With regard to gamma IFN, Torres et al (Torres, MJ, et al., Loss of an HLA haplotype in pancreas cancer tissue and its corresponding tumor derived cell line. Tissue Antigens 47:372-81, 1996) note 10 that HLA expression is upregulated by gamma IFN in pancreatic cancer, unless a total loss of haplotype has occurred. Similarly, Rees and Mian note that allelic deletion and loss can be restored, at least partially, by cytokines such as IFN-gamma (Rees, R., et al. Selective MHC expression in tumours modulates adaptive and innate antitumour responses Cancer Immunol Immunother 48:374-81, 1999). It has also been noted that IFN-gamma treatment results in upregulation of class I molecules in the majority of the cases studied 15 (Browning M, et al., Mechanisms of loss of HLA class I expression on colorectal tumor cells. Tissue Antigens 47:364-71, 1996). Kaklamakis, et al. also suggested that adjuvant immunotherapy with IFNgamma may be beneficial in the case of HLA class I negative tumors (Kaklamanis L, Loss of transporter in antigen processing 1 transport protein and major histocompatibility complex class I molecules in metastatic versus primary breast cancer. Cancer Research 55:5191-94, November 1995). It is important to underline 20 that IFN-gamma production is induced and self-amplified by local inflammation/immunization (Halloran, et al. Local T cell responses induce widespread MHC expression J. Immunol 148:3837, 1992), resulting in large increases in MHC expressions even in sites distant from the inflammatory site.

Finally, studies have demonstrated that decreased HLA expression can render tumor cells more susceptible to NK lysis (Ohnmacht, GA, et al., Heterogeneity in expression of human leukocyte 25 antigens and melanoma-associated antigens in advanced melanoma J Cellular Phys 182:332-38, 2000; Liunggren HG, et al., Host resistance directed selectively against H-2 deficient lymphoma variants: Analysis of the mechanism J. Exp. Med., 162(6):1745-59, December 1, 1985; Maio M, et al., Reduction in susceptibility to natural killer cell-mediated lysis of human FO-1 melanoma cells after induction of HLA class I antigen expression by transfection with \(\beta 2m \) gene J. Clin. Invest. 88(1):282-9, July 1991; Schrier PI, 30 et al., Relationship between myc oncogene activation and MHC class I expression Adv. Cancer Res., 60:181-246, 1993). If decreases in HLA expression benefit a tumor because it facilitates CTL escape, but render the tumor susceptible to NK lysis, then a minimal level of HLA expression that allows for resistance to NK activity would be selected for (Garrido F, et al., Implications for immunosurveillance of altered HLA class I phenotypes in human tumours Immunol Today 18(2):89-96, February 1997). Therefore, a 35 therapeutic compositions or methods in accordance with the invention together with a treatment to upregulate HLA expression and/or treatment with high affinity T-cells renders the tumor sensitive to CTL destruction.

Frequency of alterations in HLA expression

The frequency of alterations in class I expression is the subject of numerous studies (Algarra I, et al., The HLA crossroad in tumor immunology Human Immunology 61, 65-73, 2000). Rees and Mian estimate allelic loss to occur overall in 3-20% of tumors, and allelic deletion to occur in 15-50% of tumors. It should be noted that each cell carries two separate sets of class I genes, each gene carrying one HLA-A and one HLA-B locus. Thus, fully heterozygous individuals carry two different HLA-A molecules and two different HLA-B molecules. Accordingly, the actual frequency of losses for any specific allele could be as little as one quarter of the overall frequency. They also note that, in general, a gradient of expression exists between normal cells, primary tumors and tumor metastasis. In a study from Natali and coworkers (Natali PG, et al., Selective changes in expression of HLA class I polymorphic 10 determinants in human solid tumors PNAS USA 86:6719-6723, September 1989), solid tumors were investigated for total HLA expression, using W6/32 antibody, and for allele-specific expression of the A2 antigen, as evaluated by use of the BB7.2 antibody. Tumor samples were derived from primary cancers or metastasis, for 13 different tumor types, and scored as negative if less than 20%, reduced if in the 30-80% range, and normal above 80%. All tumors, both primary and metastatic, were HLA positive with W6/32. 15 In terms of A2 expression, a reduction was noted in 16.1 % of the cases, and A2 was scored as undetectable in 39.4 % of the cases. Garrido and coworkers (Garrido F, et al., Natural history of HLA expression during tumour development Immunol Today 14(10):491-99, 1993) emphasize that HLA changes appear to occur at a particular step in the progression from benign to most aggressive. Jiminez et al (Jiminez P, et al., Microsatellite instability analysis in tumors with different mechanisms for total loss of HLA expression. Cancer Immunol Immunother 48:684-90, 2000) have analyzed 118 different tumors (68 colorectal, 34 laryngeal and 16 melanomas). The frequencies reported for total loss of HLA expression were 11% for colon, 18% for melanoma and 13 % for larynx. Thus, HLA class I expression is altered in a significant fraction of the tumor types, possibly as a reflection of immune pressure, or simply a reflection of the accumulation of pathological changes and alterations in diseased cells.

Immunotherapy in the context of HLA loss

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A majority of the tumors express HLA class I, with a general tendency for the more severe alterations to be found in later stage and less differentiated tumors. This pattern is encouraging in the context of immunotherapy, especially considering that: 1) the relatively low sensitivity of 30 immunohistochemical techniques might underestimate HLA expression in tumors; 2) class I expression can be induced in tumor cells as a result of local inflammation and lymphokine release; and, 3) class I negative cells are sensitive to lysis by NK cells.

Accordingly, various embodiments of the present invention can be selected in view of the fact that there can be a degree of loss of HLA molecules, particularly in the context of neoplastic disease. For example, the treating physician can assay a patient's tumor to ascertain whether HLA is being expressed. If a percentage of tumor cells express no class I HLA, then embodiments of the present invention that comprise methods or compositions that elicit NK cell responses can be employed. As noted herein, such NK-inducing methods or composition can comprise a Flt3 ligand or ProGP which facilitate mobilization of dendritic cells, the rationale being that dendritic cells produce large amounts of IL-12. IL-

12 can also be administered directly in either amino acid or nucleic acid form. It should be noted that compositions in accordance with the invention can be administered concurrently with NK cell-inducing compositions, or these compositions can be administered sequentially.

In the context of allele-specific HLA loss, a tumor retains class I expression and may thus escape NK cell recognition, yet still be susceptible to a CTL-based vaccine in accordance with the invention which comprises epitopes corresponding to the remaining HLA type. The concept here is analogous to embodiments of the invention that include multiple disease antigens to guard against mutations that yield loss of a specific antigen. Thus, one can simultaneously target multiple HLA specificities and epitopes from multiple disease-related antigens to prevent tumor escape by allele-specific loss as well as diseaserelated antigen loss. In addition, embodiments of the present invention can be combined with alternative 10 therapeutic compositions and methods. Such alternative compositions and methods comprise, without limitation, radiation, cytotoxic pharmaceuticals, and/or compositions/methods that induce humoral antibody responses.

Moreover, it has been observed that expression of HLA can be upregulated by gamma IFN, which is commonly secreted by effector CTL, and that HLA class I expression can be induced in vivo 15 by both alpha and beta IFN. Thus, embodiments of the invention can also comprise alpha, beta and/or gamma IFN to facilitate upregualtion of HLA.

III.N. REPRIEVE PERIODS FROM THERAPIES THAT INDUCE SIDE EFFECTS: "Scheduled Treatment Interruptions or Drug Holidays"

Recent evidence has shown that certain patients infected with a pathogen, whom are initially treated with a therapeutic regimen to reduce pathogen load, have been able to maintain decreased pathogen load when removed from the therapeutic regimen, i.e., during a "drug holiday" (Rosenberg, E., et al. Immune control of HIV-1 after early treatment of acute infection Nature 407:523-26, Sept. 28, 2000) As appreciated by those skilled in the art, many therapeutic regimens for both pathogens and cancer have numerous, often severe, side effects. During the drug holiday, the patient's immune system is keeping the disease in check. Methods for using compositions of the invention are used in the context of drug holidays for cancer and pathogenic infection.

For treatment of an infection, where therapies are not particularly immunosuppressive, compositions of the invention are administered concurrently with the standard therapy. During this period, the patient's immune system is directed to induce responses against the epitopes comprised by the present inventive compositions. Upon removal from the treatment having side effects, the patient is primed to respond to the infectious pathogen should the pathogen load begin to increase. Composition of the invention can be provided during the drug holiday as well.

For patients with cancer, many therapies are immunosuppressive. Thus, upon achievement of a remission or identification that the patient is refractory to standard treatment, then upon removal from the immunosuppressive therapy, a composition in accordance with the invention is administered. Accordingly, as the patient's immune system reconstitutes, precious immune resources are simultaneously directed against the cancer. Composition of the invention can also be administered concurrently with an 40 immunosuppressive regimen if desired.

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III.O. Kits

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The peptide and nucleic acid compositions of this invention can be provided in kit form together with instructions for vaccine administration. Typically the kit would include desired peptide compositions in a container, preferably in unit dosage form and instructions for administration. An alternative kit would include a minigene construct with desired nucleic acids of the invention in a container, preferably in unit dosage form together with instructions for administration. Lymphokines such as 1L-2 or IL-12 may also be included in the kit. Other kit components that may also be desirable include, for example, a sterile syringe, booster dosages, and other desired excipients.

III.P. Overview

Epitopes in accordance with the present invention were successfully used to induce an immune response. Immune responses with these epitopes have been induced by administering the epitopes in various forms. The epitopes have been administered as peptides, as nucleic acids, and as viral vectors comprising nucleic acids that encode the epitope(s) of the invention. Upon administration of peptide-based epitope forms, immune responses have been induced by direct loading of an epitope onto an empty HLA molecule that is expressed on a cell, and via internalization of the epitope and processing via the HLA class I pathway; in either event, the HLA molecule expressing the epitope was then able to interact with and induce a CTL response. Peptides can be delivered directly or using such agents as liposomes. They can additionally be delivered using ballistic delivery, in which the peptides are typically in a crystalline form. When DNA is used to induce an immune response, it is administered either as naked DNA, generally in a dose range of approximately 1-5mg, or via the ballistic "gene gun" delivery, typically in a dose range of approximately 10-100 µg. The DNA can be delivered in a variety of conformations, e.g., linear, circular etc. Various viral vectors have also successfully been used that comprise nucleic acids which encode epitopes in accordance with the invention.

Accordingly compositions in accordance with the invention exist in several forms. Embodiments of each of these composition forms in accordance with the invention have been successfully used to induce an immune response.

One composition in accordance with the invention comprises a plurality of peptides. This plurality or cocktail of peptides is generally admixed with one or more pharmaceutically acceptable 30 excipients. The peptide cocktail can comprise multiple copies of the same peptide or can comprise a mixture of peptides. The peptides can be analogs of naturally occurring epitopes. The peptides can comprise artificial amino acids and/or chemical modifications such as addition of a surface active molecule, e.g., lipidation; acetylation, glycosylation, biotinylation, phosphorylation etc. The peptides can be CTL or HTL epitopes. In a preferred embodiment the peptide cocktail comprises a plurality of different CTL 35 epitopes and at least one HTL epitope. The HTL epitope can be naturally or non-naturally (e.g., PADRE®, Epimmune Inc., San Diego, CA). The number of distinct epitopes in an embodiment of the invention is generally a whole unit integer from one through one hundred fifty (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70,

71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100. or 150).

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An additional embodiment of a composition in accordance with the invention comprises a polypeptide multi-epitope construct, i.e., a polyepitopic peptide. Polyepitopic peptides in accordance with the invention are prepared by use of technologies well-known in the art. By use of these known technologies, epitopes in accordance with the invention are connected one to another. The polyepitopic peptides can be linear or non-linear, e.g., multivalent. These polyepitopic constructs can comprise artificial amino acids, spacing or spacer amino acids, flanking amino acids, or chemical modifications between adjacent epitope units. The polyepitopic construct can be a heteropolymer or a homopolymer. The polyepitopic constructs generally comprise epitopes in a quantity of any whole unit integer between 2-150 (e.g., 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100, or 150). The polyepitopic construct can comprise CTL and/or HTL epitopes. One or more of the epitopes in the construct can be modified, e.g., by addition of a surface active material, e.g. a lipid, or chemically modified, e.g., acetylation, etc. Moreover, bonds in the multiepitopic construct can be other than peptide bonds, e.g., covalent bonds, ester or ether bonds, disulfide bonds, hydrogen bonds, ionic bonds etc.

Alternatively, a composition in accordance with the invention comprises construct which
comprises a series, sequence, stretch, etc., of amino acids that have homology to (i.e., corresponds to or is
contiguous with to a native sequence. This stretch of amino acids comprises at least one subsequence of
amino acids that, if cleaved or isolated from the longer series of amino acids, functions as an HLA class I or
HLA class II epitope in accordance with the invention. In this embodiment, the peptide sequence is
modified, so as to become a construct as defined herein, by use of any number of techniques known or to be
provided in the art. The polyepitopic constructs can contain homology to a native sequence in any whole
unit integer increment from 70-100%, e.g., 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86,
87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, or, 100 percent.

A further embodiment of a composition in accordance with the invention is an antigen presenting cell that comprises one or more epitopes in accordance with the invention. The antigen presenting cell can be a "professional" antigen presenting cell, such as a dendritic cell. The antigen presenting cell can comprise the epitope of the invention by any means known or to be determined in the art. Such means include pulsing of dendritic cells with one or more individual epitopes or with one or more peptides that comprise multiple epitopes, by nucleic acid administration such as ballistic nucleic acid delivery or by other techniques in the art for administration of nucleic acids, including vector-based, e.g. viral vector, delivery of nucleic acids.

Further embodiments of compositions in accordance with the invention comprise nucleic acids that encode one or more peptides of the invention, or nucleic acids which encode a polyepitopic peptide in accordance with the invention. As appreciated by one of ordinary skill in the art, various nucleic acids compositions will encode the same peptide due to the redundancy of the genetic code. Each of these nucleic acid compositions falls within the scope of the present invention. This embodiment of the invention

comprises DNA or RNA, and in certain embodiments a combination of DNA and RNA. It is to be appreciated that any composition comprising nucleic acids that will encode a peptide in accordance with the invention or any other peptide based composition in accordance with the invention, falls within the scope of this invention.

5 It is to be appreciated that peptide-based forms of the invention (as well as the nucleic acids that encode them) can comprise analogs of epitopes of the invention generated using principles already known, or to be known, in the art. Principles related to analoging are now known in the art, and are disclosed herein; moreover, analoging principles (heteroclitic analoging) are disclosed in co-pending application serial number U.S.S.N. 09/226,775 filed 6 January 1999. Generally the compositions of the invention are isolated or purified.

The invention will be described in greater detail by way of specific examples. The following examples are offered for illustrative purposes, and are not intended to limit the invention in any manner. Those of skill in the art will readily recognize a variety of non-critical parameters that can be changed or modified to yield alternative embodiments in accordance with the invention.

IV. EXAMPLES

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The following example of peptide binding to HLA molecules demonstrates quantification of binding affinities of HLA class I and class II peptides. Binding assays can be performed with peptides that are either motif-bearing or not motif-bearing.

Example 1. HLA Class I and Class II Binding Assays

The following example of peptide binding to HLA molecules demonstrates quantification of binding affinities of HLA class I and class II peptides. Binding assays can be performed with peptides that are either motif-bearing or not motif-bearing.

HLA class I and class II binding assays using purified HLA molecules were performed in accordance with disclosed protocols (e.g., PCT publications WO 94/20127 and WO 94/03205; Sidney et al., Current Protocols in Immunology 18.3.1 (1998); Sidney, et al., J. Immunol. 154:247 (1995); Sette, et al., Mal. Immunol. 31:813 (1994)). Briefly, purified MHC molecules (5 to 500nM) were incubated with various unlabeled peptide inhibitors and 1-10nM ¹²³1-radiolabeled probe peptides as described. Following incubation, MHC-peptide complexes were separated from free peptide by gel filtration and the fraction of peptide bound was determined. Typically, in preliminary experiments, each MHC preparation was titered in the presence of fixed amounts of radiolabeled peptides to determine the concentration of HLA molecules necessary to bind 10-20% of the total radioactivity. All subsequent inhibition and direct binding assays were performed using these HLA concentrations.

Since under these conditions [label]- $\{HLA\}$ and $1C_{10}$ > $\{HLA\}$, the measured $1C_{10}$ values are reasonable approximations of the true K_0 values. Peptide inhibitors are typically tested at concentrations ranging from $120 \mu_2/ml$ to 1.2 ng/ml, and are tested in two to four completely independent experiments. To allow comparison of the data obtained in different experiments, a relative binding figure is calculated for each peptide by dividing the $1C_{10}$ of a positive control for inhibition by the $1C_{20}$ for each

tested peptide (typically unlabeled versions of the radiolabeled probe peptide). For database purposes, and inter-experiment comparisons, relative binding values are compiled. These values can subsequently be converted back into IC₅₀ nM values by dividing the IC₅₀ nM of the positive controls for inhibition by the relative binding of the peptide of interest. This method of data compilation has proven to be the most accurate and consistent for comparing peptides that have been tested on different days, or with different lots of ourified MHC.

Binding assays as outlined above may be used to analyze supermotif and/or motif-bearing epitopes as, for example, described in Example 2.

10 Example 2. Identification of HLA Supermotif- and Motif-Bearing CTL Candidate Epitopes

Vaccine compositions of the invention can include multiple epitopes that comprise multiple HLA supermotifs or motifs to achieve broad population coverage. This example illustrates the identification of supermotif- and motif-bearing epitopes for the inclusion in such a vaccine composition. Calculation of population coverage was performed using the strategy described below.

Computer searches and algorithms for identification of supermotif and/or motif-bearing epitopes

The searches performed to identify the motif-bearing peptide sequences in Examples 2 and 5 employed the
protein sequence data from seven proteins (E1, E2, E5, E6, E7, L1 and L2) from HPV types 16, 18, 31, 33,
45, and 56.

Accession numbers for HPV types				
Protein	6a	6b 11		
E1	Q84293	P03113	W1WL11	
2.	AAA74213	CAA25020	P04014	
		W1WL6	AAA46929	
E2	Q84294	P03119	AAA46930	
	AAA74214	CAA25021	W2WLI1	
		W2WL6	P04015	
E4	Q84295	CAA25022	P04016	
	AAA74215	W4WL6	W4WL11	
			AAA46931	
E5a	Q84296	P06460	W5WL11	
	AAA74216	CAA25023	P04017	
		W5WL6A	AAA46932	
E5b	N.A.	P06461	W5WL1B	
		CAA25024	P04018	
		W5WLB	AAA46933	
E6	O84291	P06462	W6WL11	
	AAA74211	CAA25018	P04019	
		W6WL6	AAA21703	
			AAA46927	
E7	Q84929	P06464	AAA46928	
	AAA74212	CAA25019	AAA21704	
		W7WL6	W7WL11	
			P04020	
L1	P03100	P03100	P04012	
	AAA74218	CAA25026	P1WL11	
		P1WL6	AAA4635	
L2	Q84297	P03106	P2WL11	
	1,	CAA25025	AAA46934	

		· · · · · ·	P2WL6 P040I3
	Strain	Protein Antigen	Accession number
	HPV16	E1	W1SLHS
	HPV16	E2	W2WLHS
5	HPV16	E5	W5WLHS
	HPV16	E6	W6WLHS
	HPV16	E7	W7WLHS
	HPV16	L1	AAD33259
	HPV16	L2	AAD33258
10	HPV18	E1	W1WL18
	HPV18	E2	WL18
	HPV18	E5	W5WL18
	HPV18	E6	W6WL18
	HPV18	. E7	PO6788
15	HPV18	L1	CAA28671
	HPV18	L2	P2WL18
	HPV31	E1	W1WL31
	HPV31	. E2	W2WL3
	HPV31	- E5	W5WL31
20	HPV31	E6	W6WL31
	HPV31	E7	W7WL31
	HPV31	Ll	PIWL31
	HPV31	L2	P2WL31
	HPV45	E1	S36563
25	HPV45	E2	S36564
	HPV45	E6	CAB44706
	HPV45	E7	CAB44707
	HPV45		CAB44705
	HPV45		\$36565
30	HPV33		W1WL33
	HPV33		W2WL33
	HPV33		W5WL33
	HPV33		W6WL33
	HPV33		W7WL33
35	HPV33		P1WL33
	HPV33		P2WL33
	HPV56		S36581
	HPV56		W6WL56
	HPV56		S36580
40	HPV56	L1	S38563

HPV56 L2 S36582

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Computer searches for epitopes bearing HLA Class I or Class II supermotifs or motifs were performed as follows. All translated HPV protein sequences were analyzed using a text string search software program, e.g., MotifSearch 1.4 (D. Brown, San Diego) to identify potential peptide sequences containing appropriate HLA binding motifs; alternative programs are readily produced in accordance with information in the art in view of the motif/supermotif disclosure herein. Furthermore, such calculations can be made mentally.

Identified A2., A3., and DR-supermotif sequences were scored using polynomial algorithms to predict their capacity to bind to specific HLA-Class I or Class II molecules. These polynomial algorithms take into account both extended and refined motifs (that is, to account for the impact of different amino acids at different positions), and are essentially based on the premise that the overall affinity (or ΔG) of peptide-HLA molecule interactions can be approximated as a linear polynomial function of the type:

 $\Delta G'' = a_{1i} \times a_{2i} \times a_{3i} \dots \times a_{ni}$

where a_µ is a coefficient which represents the effect of the presence of a given amino acid (f) at a given position (f) along the sequence of a peptide of a amino acids. The crucial assumption of his method is that the effects at each position are essentially independent of each other (i.e., independent binding of individual side-chains). When residue f occurs at position i in the peptide, it is assumed to contribute a constant amount f_i to the free energy of binding of the peptide irrespective of the sequence of the rest of the peptide. This assumption is justified by studies from our laboratories that demonstrated that peptides are bound to MHC and recognized by T cells in essentially an extended conformation (data omitted herein).

The method of derivation of specific algorithm coefficients has been described in

Gulukota et al., J. Mol. Biol. 267:1258-126, 1997; (see also Sidaey et al., Human Immunol. 45:79-93, 1996;
and Southwood et al., J. Immunol. 160:3363-3373, 1998). Briefly, for all i positions, anchor and nonanchor alike, the geometric mean of the average relative binding (ARB) of all peptides carrying j is
calculated relative to the remainder of the group, and used as the estimate of j. For Class II peptides, if
multiple alignments are possible, only the highest scoring alignment is utilized, following an iterative
procedure. To calculate an algorithm score of a given peptide in a test set, the ARB values corresponding
to the sequence of the peptide are multiplied. If this product exceeds a chosen threshold, the peptide is
predicted to bind. Appropriate thresholds are chosen as a function of the degree of stringency of prediction
desired.

35 Selection of HLA-A2 supertype cross-reactive peptides

Complete protein sequences from the seven HPV structural and regulatory proteins of the HPV strains listed above were aligned, then scanned, utilizing motif identification software, to identify 9and 10-mer sequences containing the HLA-A2-supermotif main anchor specificity.

HLA-A2 supermotif-bearing sequences are shown in Table VIII. Typically, these

40 sequences are then scored using the A2 algorithm and the peptides corresponding to the positive-scoring

sequences are synthesized and tested for their capacity to bind purified HLA-A*0201 molecules in vitro (HLA-A*0201 is considered a prototype A2 supertype molecule).

Examples of peptides that bind to HLA-A*0201 with IC₂₉ values <500 nM are shown in Table VIII. These peptides are then tested for the capacity to bind to additional A2-supertype molecules (A*0202, A*0203, A*0206, and A*6802). Peptides that bind to at least three of the five A2-supertype alleles tested are typically deemed A2-supertype cross-reactive binders. Preferred peptides bind at an affinity equal to or less than 500 nM to three or more HLA-A2 supertype molecules.

Selection of HLA-A3 supermotif-bearing epitopes

The HPV protein sequences scanned above were also examined for the presence of peptides with the HLA-A3-supermotif primary anchors (Table IX).

Peptides corresponding to the supermotif-bearing sequences are then synthesized and tested for binding to HLA-A*0301 and HLA-A*1101 molecules, the two most prevalent A3-supertype alleles. The peptides that are found to bind one of the two alleles with binding affinities of ≤ 500 nM, often ≤ 200 nM, are then tested for binding cross-reactivity to the other common A3-supertype alleles (A*3101, A*3301, and A*6801) to identify those that can bind at least three of the five HLA-A3-supertype molecules tested.

Selection of HLA-B7 supermotif bearing epitopes

The same HPV target antigen protein sequences were also analyzed for the presence of 9or 10-mer peptides with the HLA-B7-supermotif (Table XI).

Corresponding peptides are synthesized and tested for binding to $HLA-B^*0702$, the most common B7-supertype allele (i.e., the prototype B7 supertype allele). Peptides binding B^*0702 with IC_{50} of \$600 nM are identified using standard methods. These peptides are then tested for binding to other

25 common B7-supertype molecules (B*3501, B*5101, B*5301, and B*5401). Peptides capable of binding to three or more of the five B7-supertype alleles tested are thereby identified.

Selection of A1 and A24 motif-bearing epitopes

To further increase population coverage, HLA-A1 and -A24 epitopes can, for example,

also be incorporated into potential vaccine constructs. An analysis of the protein sequence data from the

HPV target antigens utilized above can also be performed to identify HLA-A1- and A24-motif-containing

sequences.

High affinity and/or cross-reactive binding epitopes that bear other motif and/or supermotifs are identified using analogous methodology.

Example 3. Confirmation of Immunogenicity

Cross-reactive candidate CTL A2-supermonif-bearing peptides that are identified as described in Example 2 were selected for *in vitro* immunogenicity testing. Testing was performed using the following methodology:

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Target Cell Lines for Cellular Screening:

The .221A2.1 cell line, produced by transferring the HLA-A2.1 gene into the HLA-A, -B, -C null mutant human B-lymphoblastoid cell line 721.221, is used as the peptide-loaded target to measure activity of HLA-A2.1-restricted CTL. This cell line is grown in RPMI-1640 medium supplemented with antibiotics, sodium pyruvate, nonessential amino acids and 10% (v/v) heat inactivated FCS. Cells that express an antigen of interest, or transfectants comprising the gene encoding the antigen of interest, can be used as target cells to test the ability of peptide-specific CTLs to recognize endogenous antigen.

Primary CTL Induction Cultures:

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Generation of Dendritic Cells (DC): PBMCs are thawed in RPMI with 30 g/ml DNAse, washed twice and resuspended in complete medium (RPMI-1640 plus 5% AB human serum, non-essential amino acids, sodium pyruvate, L-glutamine and penicillin/strpetomycin). The monocytes are purified by plating 10 x 10⁶ PBMC/well in a 6-well plate. After 2 hours at 37°C, the non-adherent cells are removed by gently shaking the plates and aspirating the supermatants. The wells are washed a total of three times with 3 ml RPMI to remove most of the non-adherent and loosely adherent cells. Three ml of complete medium containing 50 ng/ml of GM-CSF and 1,000 U/ml of IL-4 are then added to each well. TNF is added to the DCs on day 6 at 75 ng/ml and the cells are used for CTL induction cultures on day 7.

Induction of CTL with DC and Peptide: CD8+T-cells are isolated by positive selection with Dynal immunomagnetic beads (Dynabeads® M-450) and the detacha-bead® reagent. Typically about 200-250x10° PBMC are processed to obtain 24x10° CD8* T-cells (enough for a 48-well plate culture). Briefly, the PBMCs are thawed in RPMI with 30µg/ml DNAse, washed once with PBS containing 1% human AB serum and resuspended in PBS/1% AB serum at a concentration of 20x10° cells/ml. The magnetic beads are washed 3 times with PBS/AB serum, added to the cells (140µl beads/20x10° cells) and incubated for 1 hour at 4°C with continuous mixing. The beads and cells are washed 4x with PBS/AB serum to remove the nonadherent cells and resuspended at 100x10° cells/ml (based on the original cell number) in PBS/AB serum containing 100µl/ml detacha-bead® reagent and 30µg/ml DNAse. The mixture is incubated for 1 hour at room temperature with continuous mixing. The beads are washed again with PBS/AB/DNAse to collect the CD8+T-cells. The DC are collected and centrifuged at 1300 rpm for 5-7 minutes, washed once with PBS with 1% BSA, counted and pulsed with 40µg/ml of peptide at a cell concentration of 1-2x10°/ml in the presence of 3µg/ml B2- microglobulin for 4 hours at 20°C. The DC are then irradiated (4,200 rads), washed 1 time with medium and counted again.

Setting up induction cultures: 0.25 ml cytokine-generated DC (@1x:10⁵ cells/ml) are co-cultured with 0.25ml of CD8+T-cells (@2x10⁶ cell/ml) in each well of a 48-well plate in the presence of 10 ng/ml of IL-7. Recombinant human IL10 is added the next day at a final concentration of 10 ng/ml and rhuman IL12 is added 48 hours later at 10IU/ml.

Restimulation of the induction cultures with peptide-pulsed adherent cells: Seven and fourteen days after the primary induction the cells are restimulated with peptide-pulsed adherent cells. The PBMCS are thawed and washed twice with RPMI and DNAse. The cells are resuspended at 5x10⁶ cells'ml and irradiated at ~4200 rads. The PBMCS are plated at 2x10⁶ in 0.5ml complete medium per well and incubated for 2 hours at 37%. The plates are washed twice with RPMI by tapping the plate gently to

remove the nonadherent cells and the adherent cells pulsed with 10µg/ml of peptide in the presence of 3 µg/ml 8, microglobulin in 0.25ml RPMI/596AB per well for 2 hours at 37°C. Peptide solution from each well is aspirated and the wells are washed once with RPMI. Most of the media is aspirated from the induction cultures (CD8+ cells) and brought to 0.5 ml with fresh media. The cells are then transferred to the wells containing the peptide-pulsed adherent cells. Twenty four hours later rhuman IL10 is added at a final concentration of 10ng/ml and rhuman IL2 is added the next day and again 2-3 days later at 50IU/ml (Tsai et al., Critical Reviews in Immunology 18(1-2):65-75, 1998). Seven days later the cultures are assayed for CTL activity in a ³1Cr release assay. In some experiments the cultures are assayed for peptide-specific recognition in the in situ IFNy ELISA at the time of the second restimulation followed by assay of endogenous recognition 7 days later. After expansion, activity is measured in both assays for a side by side comparison.

Measurement of CTL lytic activity by 51Cr release.

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Seven days after the second restimulation, cytotoxicity is determined in a standard (5hr)

51Cr release assay by assaying individual wells at a single E:T. Peptide-pulsed targets are prepared by incubating the cells with 10µg/ml peptide overnight at 37°C.

Adherent target cells are removed from culture flasks with trypsin-EDTA. Target cells are labelled with 200µCi of 3 VC sodium chromate (Dupont, Wilmington, DE) for 1 hour at 37°C. Labelled target cells are resuspended at 10⁴ per ml and diluted 1:10 with K562 cells at a concentration of 3.3x10⁶/ml (an NK-sensitive erythroblastoma cell line used to reduce non-specific lysis). Target cells (100 µl) and 20 100µl of effectors are plated in 96 well round-bottom plates and incubated for 5 hours at 37°C. At that time, 100 µl of supernatant are collected from each well and percent lysis is determined according to the formula: [(cpm of the test sample- cpm of the spontaneous 3 °Cr release sample)/(cpm of the maximal 3 °Cr release sample) cpm of the spontaneous 7 °Cr release sample) (cpm of the spontaneous 7 °Cr release sample) (cpm of the spontaneous release are determined by incubating the labelled targets with 1% Trition X-100 and media alone, respectively. A positive culture is defined as one in which the specific lysis (sample- background) is 10% or higher in the case of individual wells and is 15% or more at the 2 highest E: Tratios when expanded cultures are assayed.

In situ Measurement of Human yIFN Production as an Indicator of Peptide-specific and Endogenous Recognition

Immulon 2 plates are coated with mouse anti-human IFN γ monoclonal antibody (4 µg/ml 0.1M NaHCO $_3$, pH8.2) overnight at 4°C. The plates are washed with Ca^3 , Mg^3 -free PBS/0.05% Tween 20 and blocked with PBS/10% FCS for 2 hours, after which the CTLs (100 µJ/well) and targets (100 µJ/well) are added to each well, leaving empty wells for the standards and blanks (which received media only). The target cells, either peptide-pulsed or endogenous targets, are used at a concentration of 1×10^6 cells/ml. The plates are incubated for 48 hours at 37°C with 5% CO₂.

Recombinant human IFNy is added to the standard wells starting at 400 pg or 1200pg/100µl/well and the plate incubated for 2 hours at 37°C. The plates are washed and 100 1 of biotinylated mouse anti-human IFNy monoclonal antibody (2µg/ml in PBS/3%FCS/0.05% Tween 20) are added and incubated for 2 hours at room temperature. After washing again, 100 µl HRP-streptavidin

(1:4000) are added and the plates incubated for 1 hour at room temperature. The plates are then washed 6x with wash buffer, 100µlwell developing solution (TMB 1:1) are added, and the plates allowed to develop for 5-15 minutes. The reaction is stopped with 50 µl/well 1M H₃PO₄ and read at OD450. A culture is considered positive if it measured at least 50 pg of IFNy/well above background and is twice the background level of expression.

CTL Expansion. Those cultures that demonstrate specific lytic activity against peptidepulsed targets and/or tumor targets are expanded over a two week period with anti-CD3. Briefly, 5x10⁴
CD8+ cells are added to a T25 flask containing the following: 1x10⁶ irradiated (4,200 rad) PBMC
dutologous or allogencic) per ml, 2x10⁵ irradiated (8,000 rad) EBV- transformed cells per ml, and OKT3
(anti-CD3) at 30ng per ml in RPMI-1640 containing 10% (v/v) human AB serum, non-essential amino
acids, sodium pytuvate, 25µM 2-mercaptoethanol, L-glutamine and penicillin/streptomycin. Rhuman IL2
is added 24 hours later at a final concentration of 200IU/ml and every 3 days thereafter with fresh media at
50IU/ml. The cells are split if the cell concentration exceeded 1x10⁶/ml and the cultures are assayed
between days 13 and 15 at E:T ratios of 30, 10, 3 and 1:1 in the ¹⁶Cr release assay or at 1x10⁶/ml in the in
stru IFN assay using the same targets as before the expansion.

Cultures are expanded in the absence of anti-CD3* as follows. Those cultures that demonstrate specific lytic activity against peptide and endogenous targets are selected and Sx10* CD8* cells are added to a T25 flask containing the following: 1x10* autologous PBMC per ml which have been peptide-pulsed with 10µg/ml peptide for 2 hours at 37°C and irradiated (4,200 rad); 2x10* irradiated (8,000 rad) EBV-transformed cells per ml RPMI-1640 containing 10%(v/v) human AB serum, non-essential AA, sodium pyruvate, 25mM 2-ME, L-glutamine and gentamicin.

Immunogenicity of A2 supermotif-bearing peptides

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A2-supermotif cross-reactive binding peptides are tested in the cellular assay for the

ability to induce peptide-specific CTL in normal individuals. In this analysis, a peptide is typically
considered to be an epitope if it induces peptide-specific CTLs in at least 2 donors (unless otherwise noted)
and preferably, also recognizes the endogenously expressed peptide.

Immunogenicity is additionally confirmed using PBMCs isolated from HPV-infected patients. Briefly, PBMCs are isolated from patients, re-stimulated with peptide-pulsed monocytes and assayed for the ability to recognize peptide-pulsed target cells as well as transfected cells endogenously expressing the antigen.

Evaluation of A*03/A11 immunogenicity

HLA-A3 supermotif-bearing cross-reactive binding peptides are also evaluated for
immunogenicity using methodology analogous for that used to evaluate the immunogenicity of the HLA-A2 supermotif peptides.

Evaluation of B7 immunogenicity

Immunogenicity screening of the B7-supertype cross-reactive binding peptides identified in Example 2 are evaluated in a manner analogous to the evaluation of A2-and A3-supermotif-bearing peptides.

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Peptides bearing other supermotifs/motifs, e.g., HLA-A1, HLA-A24 etc. are also evaluated using similar methodology

Example 4. Implementation of the Extended Supermotif to Improve the Binding Capacity of Native Epitopes by Creating Analogs

HLA motifs and supermotifs (comprising primary and/or secondary residues) are useful in the identification and preparation of highly cross-reactive native peptides, as demonstrated herein. Moreover, the definition of HLA motifs and supermotifs also allows one to engineer highly cross-reactive epitopes by identifying residues within a native peptide sequence which can be analoged, or "fixed" to confer upon the peptide certain characteristics, e.g. greater cross-reactivity within the group of HLA molecules that comprise a supertype, and/or greater binding affinity for some or all of those HLA molecules. Examples of analoging peptides to exhibit modulated binding affinity are set forth in this example.

20 Analoging at Primary Anchor Residues

Peptide engineering strategies are implemented to further increase the cross-reactivity of the epitopes. For example, on the basis of the data disclosed, e.g., in related and co-pending U.S.S.N 09/226,775, the main anchors of A2-supermotif-bearing peptides are altered, for example, to introduce a preferred L, I, V, or M at position 2, and I or V at the C-terminus.

To analyze the cross-reactivity of the analog peptides, each engineered analog is initially tested for binding to the prototype A2 supertype allele A*0201, then, if A*0201 binding capacity is maintained, for A2-supertype cross-reactivity.

Alternatively, a peptide is tested for binding to one or all supertype members and then analogued to modulate binding affinity to any one (or more) of the supertype members to add population coverage.

The selection of analogs for immunogenicity in a cellular screening analysis is typically further restricted by the capacity of the parent peptide to bind at least weakly, i.e., bind at an IC₅₀ of 5000nM or less, to three of more A2 supertype alleles. The rationale for this requirement is that the WT peptides must be present endogenously in sufficient quantity to be biologically relevant. Analoged peptides have been shown to have increased immunogenicity and cross-reactivity by T cells specific for the parent epitope (see, e.g., Parkhurst et al., J. Immunol. 157:2539, 1996; and Pogue et al., Proc. Natl. Acad. Sci. USA 92:8166, 1995).

In the cellular screening of these peptide analogs, it is important to demonstrate that analog-specific CTLs are also able to recognize the wild-type peptide and, when possible, target cells that 40 endogenously express the epitope.

Analoging of HLA-A3 and B7-supermotif-bearing peptides

Analogs of HLA-A3 supermotif-bearing epitopes are generated using strategies similar to those employed in analoging HLA-A2 supermotif-bearing peptides. For example, peptides binding to 3/5 of the A3-supertype molecules are engineered at primary anchor residues to possess a preferred residue (V, S, M, or A) at position 2.

The analog peptides are then tested for the ability to bind A^*03 and A^*11 (prototype A3 supertype alleles). Those peptides that demonstrate ≤ 500 nM binding capacity are then tested for A3-supertype cross-reactivity.

Similarly to the A2- and A3- motif bearing peptides, peptides binding 3 or more B7-supertype alleles can be improved, where possible, to achieve increased cross-reactive binding. B7 supermotif-bearing peptides are, for example, engineered to possess a preferred residue (V, I, L, or F) at the C-terminal primary anchor position, as demonstrated by Sidney et al. (J. Immunol. 157:3480-3490, 1996).

Analoguing at primary anchor residues of other motif and/or supermotif-bearing epitopes is performed in a like manner.

The analog peptides are then be tested for immunogenicity, typically in a cellular screening assay. Again, it is generally important to demonstrate that analog-specific CTLs are also able to recognize the wild-type peptide and, when possible, targets that endogenously express the epitope.

20 Analoging at Secondary Anchor Residues

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Moreover, HLA supermotifs are of value in engineering highly cross-reactive peptides and/or peptides that bind HLA molecules with increased affinity by identifying particular residues at secondary anchor positions that are associated with such properties. For example, the binding capacity of a B7 supermotif-bearing peptide with an F residue at postion 1 is analyzed. The peptide is then analoged to, for example, substitute L for F at position 1. The analoged peptide is evaluated for increased binding affinity/ and or increased cross-reactivity. Such a procedure identifies analoged peptides with modulated binding affinity.

Engineered analogs with sufficiently improved binding capacity or cross-reactivity can also be tested for immunogenicity in HLA-B7-transgenic mice, following for example, IFA immunization or lipopeptide immunization. Analogued peptides are additionally tested for the ability to stimulate a recall response using PBMC from HPV-infected patients.

Other analoguing strategies

Another form of peptide analoguing, unrelated to the anchor positions, involves the substitution of a cysteine with α-amino butyric acid. Due to its chemical nature, cysteine has the propensity to form disulfide bridges and sufficiently after the peptide structurally so as to reduce binding capacity. Subtitution of α-amino butyric acid for cysteine not only alleviates this problem, but has been shown to improve binding and crossbinding capabilities in some instances (see, e.g., the review by Sette et al., In: Persistent Viral Infections, Eds. R. Ahmed and I. Chen, John Wiley & Sons, England, 1999).

Thus, by the use of even single amino acid substitutions, the binding affinity and/or crossreactivity of peptide ligands for HLA supertype molecules can be modulated.

Example 5. Identification of HPV-derived sequences with HLA-DR binding motifs

Peptide epitopes bearing an HLA class II supermotif or motif are identified as outlined below using methodology similar to that described in Examples 1-3.

Selection of HLA-DR-supermotif-bearing epitopes.

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To identify HPV-derived, HLA class II HTL epitopes, the protein sequences from the same HPV antigens used for the identification of HLA Class I supermotif/motif sequences were analyzed for the presence of sequences bearing an HLA-DR-motif or supermotif. Specifically, 15-mer sequences were selected comprising a DR-supermotif, further comprising a 9-mer core, and three-residue N- and Cterminal flanking regions (15 amino acids total).

Protocols for predicting peptide binding to DR molecules have been developed

(Southwood et al., J. Immunol. 160:3363-3373, 1998). These protocols, specific for individual DR molecules, allow the scoring, and ranking, of 9-mer core regions. Each protocol not only scores peptide sequences for the presence of DR-supermotif primary anchors (i.e., at position 1 and position 6) within a 9-mer core, but additionally evaluates sequences for the presence of secondary anchors. Using allele specific selection tables (see, e.g., Southwood et al., ibid.), it has been found that these protocols efficiently select 20 peptide sequences with a high probability of binding a particular DR molecule. Additionally, it has been found that performing these protocols in tandem, specifically those for DR1, DR4w4, and DR7, can efficiently select DR cross-reactive peptides.

The HPV-derived peptides identified above are tested for their binding capacity for various common HLA-DR molecules. All peptides are initially tested for binding to the DR molecules in the primary panel: DR1, DR4w4, and DR7. Peptides binding at least 2 of these 3 DR molecules are then tested for binding to DR2w2 β1, DR2w2 β2, DR6w19, and DR9 molecules in secondary assays. Finally, peptides binding at least 2 of the 4 secondary panel DR molecules, and thus cumulatively at least 4 of 7 different DR molecules, are screened for binding to DR4w15, DR5w11, and DR8w2 molecules in tertiary assays. Peptides binding at least 7 of the 10 DR molecules comprising the primary, secondary, and tertiary screening assays are considered cross-reactive DR binders. HPV-derived peptides found to bind common HLA-DR alleles are of particular interest.

Selection of DR3 motif peptides

Because HLA-DR3 is an allele that is prevalent in Caucasian, Black, and Hispanic populations, DR3 binding capacity is an important criterion in the selection of HTL epitopes. However, data generated previously indicated that DR3 only rarely cross-reacts with other DR alleles (Sidney et al., J. Immunol. 149:2634-2640, 1992; Geluk et al., J. Immunol. 152:5742-5748, 1994; Southwood et al., J. Immunol. 160:3363-3373, 1998). This is not entirely surprising in that the DR3 peptide-binding motif appears to be distinct from the specificity of most other DR alleles. For maximum efficiency in developing vaccine candidates it would be desirable for DR3 motifs to be clustered in proximity with DR supermotif

regions. Thus, peptides shown to be candidates may also be assayed for their DR3 binding capacity.

However, in view of the distinct binding specificity of the DR3 motif, peptides binding only to DR3 can also be considered as candidates for inclusion in a vaccine formulation.

To efficiently identify peptides that bind DR3, target HPV antigens are analyzed for sequences carrying one of the two DR3 specific binding motifs reported by Geluk et al. (1. Immunol. 152:5742-5748, 1994). The corresponding peptides are then synthesized and tested for the ability to bind DR3 with an affinity of 1µM or better, i.e., less than 1 µM. Peptides are found that meet this binding criterion and qualify as HLA class II high affinity binders.

DR3 binding epitopes identified in this manner are included in vaccine compositions with

10 DR supermotif-bearing peptide epitopes.

Similarly to the case of HLA class I motif-bearing peptides, the class II motif-bearing peptides are analoged to improve affinity or cross-reactivity. For example, aspartic acid at position 4 of the 9-mer core sequence is an optimal residue for DR3 binding, and substitution for that residue often improves DR 3 binding.

Example 6. Immunogenicity of HPV-derived HTL epitopes

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This example determines immunogenic DR supermotif- and DR3 motif-bearing epitopes among those identified using the methodology in Example 5.

Immunogenicity of HTL epitopes are evaluated in a manner analogous to the 20 determination of immunogenicity of CTL epitopes by assessing the ability to stimulate HTL responses and/or by using appropriate transgenic mouse models. Immunogenicity is determined by screening for: 1.) in viro primary induction using normal PBMC or 2.) recall responses from cancer patient PBMCs.

Example 7. Calculation of phenotypic frequencies of HLA-supertypes in various ethnic backgrounds to determine breadth of population coverage

This example illustrates the assessment of the breadth of population coverage of a vaccine composition comprised of multiple epitopes comprising multiple supermotifs and/or motifs.

In order to analyze population coverage, gene frequencies of HLA alleles were determined. Gene frequencies for each HLA allele were calculated from antigen or allele frequencies utilizing the binomial distribution formulae gf=1-{SQRT(1-af)} (see, e.g., Sidney et al., Human Immunol. 45:79-93, 1996). To obtain overall phenotypic frequencies, cumulative gene frequencies were calculated, and the cumulative antigen frequencies derived by the use of the inverse formula [af=1-(1-Cgf)²].

Where frequency data was not available at the level of DNA typing, correspondence to the serologically defined antigen frequencies was assumed. To obtain total potential supertype population coverage no linkage disequilibrium was assumed, and only alleles confirmed to belong to each of the supertypes were included (minimal estimates). Estimates of total potential coverage achieved by inter-loci combinations were made by adding to the A coverage the proportion of the non-A covered population that could be expected to be covered by the B alleles considered (e.g., total=A+B*q1-A)). Confirmed members of the A3-like supertype are A3, A11, A31, A3301, and A*6801. Although the A3-like supertype may also include A34, A66, and A*7401, these alleles were not included in overall frequency calculations.

Likewise, confirmed members of the A2-like supertype family are A*0201, A*0202, A*0203, A*0204, A*0205, A*0206, A*0207, A*6802, and A*6901. Finally, the B7-like supertype-confirmed alleles are: B7, B*3501-03, B51, B*5301, B*5401, B*5501-2, B*5601-2, B*6701, and B*7801 (potentially also B*1401, B*3504-06, B*4201, and B*5602).

Population coverage achieved by combining the A2-, A3- and B7-supertypes is approximately 86% in five major ethnic groups, supra. Coverage may be extended by including peptides bearing the A1 and A24 motifs. On average, A1 is present in 12% and A24 in 29% of the population across five different major ethnic groups (Caucasian, North American Black, Chinese, Japanese, and Hispanic). Together, these alleles are represented with an average frequency of 39% in these same ethnic populations. The total coverage across the major ethnicities when A1 and A24 are combined with the coverage of the A2-, A3- and B7-supertype alleles is >95%. An analagous approach can be used to estimate population coverage achieved with combinations of class II motif-bearing epitopes.

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Immunogenicity studies in humans (e.g., Bertoni et al., J. Clin. Invest. 100.503, 1997; Doolan et al., Immunity 7:97, 1997; and Threlkeld et al., J. Immunol. 159:1648, 1997) have shown that highly cross-reactive hinding peptides are almost always recognized as epitopes. The use of highly cross-reactive binding peptides is an important selection criterion in identifying candidate epitopes for inclusion in a vaccine that is immunogenic in a diverse population.

With a sufficient number of epitopes (as disclosed herein and from the art), an average population coverage is predicted to be greater than 95% in each of five major ethnic populations. The game theory Monte Carlo simulation analysis, which is known in the art (see e.g., Osborne, M.J. and Rubinstein, A. "A course in game theory" MIT Press, 1994), can be used to estimate what percentage of the individuals in a population comprised of the Caucasian, North American Black, Japanese, Chinese, and Hispanic ethnic groups would recognize the vaccine epitopes described herein. A preferred percentage is 90%. A more preferred percentage is 95%.

Example 8. CTL Recognition Of Endogenous Processed Antigens After Priming

This example determines that CTL induced by native or analogued peptide epitopes identified and selected as described in Examples 1-6 recognize endogenously synthesized, i.e., native antigens.

Effector cells isolated from transgenic mice that are immunized with peptide epitopes as in Example 3, for example HLA-A2 supermotif-bearing epitopes, are re-stimulated in viron using peptide-coated stimulator cells. Six days later, effector cells are assayed for cytotoxicity and the cell lines that contain peptide-specific cytotoxic activity are further re-stimulated. An additional six days later, these cell ines are tested for cytotoxic activity on 51 Cr labeled Jurkat-A2. $1/K^{5}$ target cells in the absence or presence of peptide, and also tested on 51 Cr labeled target cells bearing the endogenously synthesized antigen, i.e. cells that are stably transfected with HPV expression vectors.

The result will demonstrate that CTL lines obtained from animals primed with peptide epitope recognize endogenously synthesized HPV antigen. The choice of transgenic mouse model to be used for such an analysis depends upon the epitope(s) that is being evaluated. In addition to HLA-A **O201/K** transgenic mice, several other transgenic mouse models including mice with human A11, which

may also be used to evaluate A3 epitopes, and B7 alleles have been characterized and others (e.g., transgenic mice for HLA-A1 and A24) are being developed. HLA-DR1 and HLA-DR3 mouse models have also been developed, which may be used to evaluate HTL epitopes.

5 Example 9. Activity Of CTL-HTL Conjugated Epitopes In Transgenic Mice

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This example illustrates the induction of CTLs and HTLs in transgenic mice by use of a tumor associated antigen CTL/HTL peptide conjugate whereby the vaccine composition comprises peptides to be administered to an HPV-infected patient. The peptide composition can comprise multiple CTL and/or HTL epitopes and further, can comprise epitopes selected from multiple HPV target antigens. The epitopes are identified using methodology as described in Examples 1-6 This analysis demonstrates the enhanced immunogenicity that can be achieved by inclusion of one or more HTL epitopes in a vaccine composition. Such a peptide composition can comprise an HTL epitope conjugated to a preferred CTL epitope containing, for example, at least one CTL epitope that binds to multiple HLA family members at an affinity of 500 nM or less, or analogs of that epitope. The peptides may be lipidated, if desired.

Immunization procedures: Immunization of transgenic mice is performed as described (Alexander et al., J. Immunol. 159:4753-4761, 1997). For example, A2/K³ mice, which are transgenic for the human HLA A2.1 allele and are useful for the assessment of the immunogenicity of HLA-A*02.01 motif- or HLA-A*2 supermotif-bearing epitopes, are primed subcutaneously (base of the tail) with a 0.1 ml of peptide in Incomplete Freund's Adjuvant, or if the peptide composition is a lipidated CTL/HTL conjugate, in DMSO/saline or if the peptide composition is a polypeptide, in PBS or Incomplete Freund's Adjuvant. Seven days after priming, splenocytes obtained from these animals are restimulated with synaenic irradiated LPS-activated lymphoblasts coated with peptide.

Cell lines: Target cells for peptide-specific cytotoxicity assays are Jurkat cells transfected with the HLA-A2.1/Kb chimeric gene (e.g., Vitiello et al., J. Exp. Med. 173:1007, 1991)

In vitro CTL activation: One week after priming, spleen cells (30x10⁶ cells/flask) are cocultured at 37°C with syngeneic, irradiated (3000 rads), peptide coated lymphoblasts (10x10⁶ cells/flask) in 10 ml of culture medium/T25 flask. After six days, effector cells are harvested and assayed for cytotoxic activity.

Assay for cytotoxic activity: Target cells $(1.0 \text{ to } 1.5 \times 10^6)$ are incubated at 37^8C in the presence of $200 \, \mu \text{l}$ of ^{51}Cr . After 60 minutes, cells are washed three times and resuspended in R10 medium. Peptide is added where required at a concentration of 1 $\mu \text{g/ml}$. For the assay, 10^6 ^{51}Cr -labeled target cells are added to different concentrations of effector cells (final volume of $200 \, \mu \text{l}$) in U-bottom 96-well plates. After a 6 hour incubation period at 37^8C , a 0.1 ml aliquot of supernatant is removed from each well and radioactivity is determined in a Micromedic automatic gamma counter. The percent specific lysis is determined by the formula: percent specific release = $100 \times (\text{experimental release} - \text{spontaneous}$ release)/(maximum release - spontaneous release). To facilitate comparison between separate CTL assays run under the same conditions, $96 \, ^{10}\text{Cr}$ release data is expressed as lytic units/ 10^6 cells. One lytic unit is arbitrarily defined as the number of effector cells required to achieve 30% lysis of 10,000 target cells in a 6 hour ^{11}Cr release assay. To obtain specific lytic units/ 10^6 , the lytic units/ 10^6 obtained in the absence of peptide. For example, if $30\% \, ^{11}\text{Cr}$

release is obtained at the effector (E): target (T) ratio of 50:1 (i.e., 5×10^5 effector cells for 10,000 targets) in the absence of peptide and 5:1 (i.e., 5×10^5 effector cells for 10,000 targets) in the presence of peptide, the specific lytic units would be: $[(1/50,000) \cdot (1/500,000)] \times 10^6 = 18$ LU.

The results are analyzed to assess the magnitude of the CTL responses of animals injected with the immunogenic CTL/HTL conjugate vaccine preparation and are compared to the magnitude of the CTL response achieved using the CTL epitope as outlined in Example 3. Analyses similar to this may be performed to evaluate the immunogenicity of peptide conjugates containing multiple CTL epitopes and/or multiple HTL epitopes. In accordance with these procedures it is found that a CTL response is induced, and concominantly that an HTL response is induced upon administration of such compositions.

Example 10. Selection of CTL and HTL epitopes for inclusion in an HPV-specific vaccine.

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This example illustrates the procedure for the selection of peptide epitones for vaccine compositions of the invention. The peptides in the composition can be in the form of a nucleic acid sequence, either single or one or more sequences (i.e., minigene) that encodes peptide(s), or can be single and/or nolveolitonic peptides.

The following principles are utilized when selecting an array of epitopes for inclusion in a vaccine composition. Each of the following principles is balanced in order to make the selection.

Epitopes are selected which, upon administration, mimic immune responses that have been observed to be correlated with HPV clearance. The number of épitopes used depends on observations of patients who spontaneously clear HPV. For example, if it has been observed that patients who spontaneously clear HPV generate an immune response to at least 3 epitopes on at least one HPV antigen, then 3-4 epitopes should be included for HLA class I. A similar rationale is used to determine HLA class II epitopes.

When selecting an array of HPV epitopes, it is preferred that at least some of the epitopes are derived from early and late proteins. The early proteins of HPV are expressed when the virus is replicating, either following acute or dormant infection. Therefore, it is particularly preferred to use epitopes from early stage proteins to alleviate disease manifestations at the earliest stage possible.

Epitopes are often selected that have a binding affinity of an IC_{50} of 500 nM or less for an HLA class I molecule, or for class II, an IC_{50} of 1000 nM or less.

Sufficient supermotif bearing peptides, or a sufficient array of allele-specific motif bearing peptides, are selected to give broad population coverage. For example, epitopes are selected to provide at least 80% population coverage. A Monte Carlo analysis, a statistical evaluation known in the art, can be employed to assess breadth, or redundancy, of population coverage.

When creating a polyepitopic compositions, e.g. a minigene, it is typically desirable to generate the smallest peptide possible that encompasses the epitopes of interest. The principles employed are similar, if not the same, as those employed when selecting a peptide comprising nested epitopes.

In cases where the sequences of multiple variants of the same target protein are available, potential peptide epitopes can also be selected on the basis of their conservancy. For example, a criterion for conservancy may define that the entire sequence of an HLA class I binding peptide or the entire 9-mer

core of a class II binding peptide be conserved in a designated percentage of the sequences evaluated for a specific protein antigen.

A vaccine composition comprised of selected peptides, when administered, is safe, efficacious, and elicits an immune response similar in magnitude to an immune response that controls or clears an acute HPV infection.

Example 11. Construction of Minigene Multi-Epitope DNA Plasmids

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This example provides general guidance for the construction of a minigene expression plasmid. Minigene plasmids may, of course, contain various configurations of CTL and/or HTL epitopes or 10 epitope analogs as described herein. Examples of the construction and evaluation of expression plasmids are described, for example, in co-pending U.S.S.N. 09/311,784 filed 5/13/99.

A minigene expression plasmid typically includes multiple CTL and HTL peptide epitopes. In the present example, HLA-A2, -A3, -B7 supermotif-bearing peptide epitopes and HLA-A1 and -A24 motif-bearing peptide epitopes are used in conjunction with DR supermotif-bearing peptide epitopes and HLA-A1 and -A24 motif-bearing peptide epitopes. HLA class I supermotif or motif-bearing peptide epitopes derived from multiple HPV antigens, preferably including both early and late phase antigens, are selected such that multiple supermotifs/motifs are represented to ensure broad population coverage. Similarly, HLA class II epitopes are selected from multiple HPV antigens to provide broad population coverage, t.e. both HLA DR-1-4-7 supermotif-bearing epitopes and HLA DR-3 motif-bearing epitopes are selected for inclusion in the minigene construct. The selected CTL and HTL epitopes are then incorporated into a minigene for expression in an expression vector.

Such a construct may additionally include sequences that direct the HTL epitopes to the endoplasmic reticulum. For example, the II protein may be fused to one or more HTL epitopes as described in co-pending application U.S.S.N. 09/311,784 filed 5/13/99, wherein the CLIP sequence of the II protein is removed and replaced with an HLA class II epitope sequence so that HLA class II epitope is directed to the endoplasmic reticulum, where the epitope binds to an HLA class II molecules.

This example illustrates the methods to be used for construction of a minigene-bearing expression plasmid. Other expression vectors that may be used for minigene compositions are available and known to those of skill in the art.

The minigene DNA plasmid of this example contains a consensus Kozak sequence and a consensus murine kappa Ig-light chain signal sequence followed by CTL and/or HTL epitopes selected in accordance with principles disclosed herein. The sequence encodes an open reading frame fused to the Myc and His antibody epitope tag coded for by the pcDNA 3.1 Myc-His vector.

Overlapping oligonucleotides that can, for example, average about 70 nucleotides in length with 15 nucleotide overlaps, are synthesized and HPLC-purified. The oligonucleotides encode the selected peptide epitopes as well as appropriate linker nucleotides, Kozak sequence, and signal sequence. The final multiepitope minigene is assembled by extending the overlapping oligonucleotides in three sets of reactions using PCR. A Perkin/Elmer 9600 PCR machine is used and a total of 30 cycles are performed using the following conditions: 95°C for 15 sec, annealing temperature (5° below the lowest calculated Tm of each primer pair) for 30 sec, and 72°C for 1 min.

For example, a minigene can be prepared as follows. For a first PCR reaction, $5 \mu g$ of each of two oligonucleotides are annealed and extended: In an example using eight oligonucleotides, i.e., four pairs of primers, oligonucleotides 1+2, 3+4, 5+6, and 7+8 are combined in 100 μ 1 reactions containing Pfu polymerase buffer (1x=10 mM KCL, 10 mM (NH4)₂SO₄, 20 mM Tris-chloride, pH 8.75, 2 mM MSO₄, 0.1% Triton X-100, 100 μ 2/ml BSA), 0.25 mM each dNTP, and 2.5 U of Pfu polymerase. The full-length dimer products are gel-purified, and two reactions containing the product of 1+2 and 3+4, and the product of 5+6 and 7+8 are mixed, annealed, and extended for 10 cycles. Half of the two reactions are then mixed, and 5 cycles of annealing and extension carried out before flanking primers are added to amplify the full length product. The full-length product is gel-purified and cloned into pCR-blunt (Invirogen) and individual clones are screened by sequencing.

Example 12. The plasmid construct and the degree to which it induces immunogenicity.

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The degree to which a plasmid construct, for example a plasmid constructed in accordance with Example 11, is able to induce immunogenicity can be evaluated in vitro by testing for epitope presentation by APC following transduction or transfection of the APC with an epitope-expressing nucleic acid construct. Such a study determines "antigenicity" and allows the use of human APC. The assay determines the ability of the epitope to be presented by the APC in a context that is recognized by a T cell by quantifying the density of epitope-HLA class I complexes on the cell surface. Quantitation can be performed by directly measuring the amount of peptide eluted from the APC (see, e.g., Sijts et al., J. Immunol. 136:683-692, 1996; Demotz et al., Nature 342:682-684, 1989); or the number of peptide-HLA class I complexes can be estimated by measuring the amount of lysis or lymphokine release induced by infected or transfected target cells, and then determining the concentration of peptide necessary to obtained equivalent levels of lysis or lymphokine release (see, e.g., Kageyama et al., J. Immunol. 154:567-576, 1995).

Atternatively, immunogenicity can be evaluated through in vivo injections into mice and subsequent in vitro assessment of CTL and HTL activity, which are analysed using cytotoxicity and proliferation assays, respectively, as detailed e.g., in copending U.S.S.N. 09/311,784 filed 5/13/99 and Alexander et al., Immunity 1:751-761, 1994.

For example, to assess the capacity of a DNA minigene construct (e.g., a pMin minigene construct generated as decribed in U.S.S.N. 09/311/784) containing at least one HLA-A2 supermotif peptide to induce CTLs in vivo, HLA-A2.1/K^b transgenic mice, for example, are immunized intramuscularly with 100 µg of naked cDNA. As a means of comparing the level of CTLs induced by CDNA immunization, a control group of animals is also immunized with an actual peptide composition that comprises multiple epitopes synthesized as a single polypeptide as they would be encoded by the minigene.

Splenocytes from immunized animals are stimulated twice with each of the respective compositions (peptide epitopes encoded in the minigene or the polyepitopic peptide), then assayed for peptide-specific cytotoxic activity in a ³¹Cr release assay. The results indicate the magnitude of the CTL response directed against the A2-restricted epitope, thus indicating the *in vivo* immunogenicity of the minigene vaccine and polyepitopic vaccine. It is, therefore, found that the minigene elicits immune responses directed toward the HLA-A2 supermotif peptide epitopes as does the polyepitopic peptide

vaccine. A similar analysis is also performed using other HLA-A3 and HLA-B7 transgenic mouse models to assess CTL induction by HLA-A3 and HLA-B7 motif or supermotif epitopes.

To assess the capacity of a class II epitope encoding minigene to induce HTLs in wwo, DR transgenic mice, or for those epitope that cross react with the appropriate mouse MHC molecule, I-A²restricted mice, for example, are immunized intramuscularly with 100 µg of plasmid DNA. As a means of comparing the level of HTLs induced by DNA immunization, a group of control animals is also immunized with an actual peptide composition emulsified in complete Freund's adjuvant. CD4+ T cells, i.e. HTLs, are purified from splenocytes of immunized animals and stimulated with each of the respective compositions (peptides encoded in the minigene). The HTL response is measured using a ³H-thymidine incorporation proliferation assay, (see, e.g., Alexander et al. Immunity 1:751-761, 1994). The results indicate the magnitude of the HTL response, thus demonstrating the in vivo immunogenicity of the minigene.

DNA minigenes, constructed as described in Example 11, may also be evaluated as a vaccine in combination with a boosting agent using a prime boost protocol. The boosting agent can consist of recombinant protein (e.g., Bamett et al., Aids Res. and Human Retroviruses 14, Supplement 3:S299-S309, 1998) or recombinant vaccinia, for example, expressing a minigene or DNA encoding the complete protein of interest (see, e.g., Hanke et al., Vaccine 16:439-445, 1998; Sedegah et al., Proc. Natl. Acad. Sci USA 95:7648-53, 1998; Hanke and McMichael, Immunol. Letters 66:177-181, 1999; and Robinson et al., Nature Med. 5:526-34, 1999).

For example, the efficacy of the DNA minigene used in a prime boost protocol is initially

20 evaluated in transgenic mice. In this example, A2.1/K* transgenic mice are immunized IM with 100 g of

a DNA minigene encoding the immunogenic peptides including at least one HLA-A2 supermotif-bearing
peptide. After an incubation period (ranging from 3-9 weeks), the mice are boosted IP with 10' pfu/mouse
of a recombinant vaccinia virus expressing the same sequence encoded by the DNA minigene. Control
mice are immunized with 100 g of DNA or recombinant vaccinia without the minigene sequence, or with
DNA encoding the minigene, but without the vaccinia boost. After an additional incubation period of two
weeks, splenocytes from the mice are immediately assayed for peptide-specific activity in an ELISPOT
assay. Additionally, splenocytes are stimulated in viro with the A2-restricted peptide epitopes encoded in
the minigene and recombinant vaccinia, then assayed for peptide-specific activity in an IFNELISA.

It is found that the minigene utilized in a prime-boost protocol elicits greater immune
responses toward the HLA-A2 supermotif peptides than with DNA alone. Such an analysis can also be
performed using HLA-A11 or HLA-B7 transgenic mouse models to assess CTL induction by HLA-A3 or
HLA-B7 motif or supermotif epitopes.

The use of prime boost protocols in humans is described in Example 20.

35 Example 13. Peptide Composition for Prophylactic Uses

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Vaccine compositions of the present invention can be used to prevent HPV infection in persons who are at risk for such infection. For example, a polyepitopic peptide epitope composition (or a nucleic acid comprising the same) containing multiple CTL and HTL epitopes such as those selected in Examples 9 and/or 10, which are also selected to target greater than 80% of the population, is administered to individuals at risk for HPV infection.

For example, a peptide-based composition can be provided as a single polypeptide that encompasses multiple epitopes. The vaccine is typically administered in a physiological solution that comprises an adjuvant, such as Incomplete Freunds Adjuvant. The dose of peptide for the initial immunization is from about 1 to about 50,000 µg, generally 100-5,000 µg, for a 70 kg patient. The initial administration of vaccine is followed by booster dosages at 4 weeks followed by evaluation of the magnitude of the immune response in the patient, by techniques that determine the presence of epitope-specific CTL populations in a PBMC sample. Additional booster doses are administrated as required. The composition is found to be both safe and efficacious as a prophylaxis against HPV infection.

Alternatively, a composition typically comprising transfecting agents can be used for the

administration of a nucleic acid-based vaccine in accordance with methodologies known in the art and
disclosed herein

Example 14. Polyepitopic Vaccine Compositions Derived from Native HPV Sequences

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A native HPV polyprotein sequence is screened, preferably using computer algorithms defined for each class I and/or class II supermotif or motif, to identify "relatively short" regions of the polyprotein that comprise multiple epitopes and is preferably less in length than an entire native antigen. This relatively short sequence that contains multiple distinct, even overlapping, epitopes is selected and used to generate a minigene construct. The construct is engineered to express the peptide, which corresponds to the native protein sequence. The "relatively short" peptide is generally less than 250 amino acids in length, often less than 100 amino acids in length, preferably less than 75 amino acids in length, and more preferably less than 50 amino acids in length. The protein sequence of the vaccine composition is selected because it has maximal number of epitopes contained within the sequence, i.e., it has a high concentration of epitopes. As noted herein, epitope motifs may be nested or overlapping (i.e., frame shifted relative to one another). For example, with foverlapping epitopes, two 9-mer epitopes and one 10-mer epitope can be present in a 10 amino acid peptide. Such a vaccine composition is administered for therapeutic or prophylactic purposes.

The vaccine composition will include, for example, three CTL epitopes from at least one HPV target antigen and at least one HTL epitope. This polyepitopic native sequence is administered either as a peptide or as a nucleic acid sequence which encodes the peptide. Alternatively, an analog can be made of this native sequence, whereby one or more of the epitopes comprise substitutions that alter the cross-reactivity and/or binding affinity properties of the polyepitopic peptide.

The embodiment of this example provides for the possibility that an as yet undiscovered aspect of immune system processing will apply to the native nested sequence and thereby facilitate the production of therapeutic or prophylactic immune response-inducing vaccine compositions. Additionally such an embodiment provides for the possibility of motif-bearing epitopes for an HLA makeup that is presently unknown. Furthermore, this embodiment (absent analogs) directs the immune response to multiple peptide sequences that are actually present in native HPV antigens thus avoiding the need to evaluate any junctional epitopes. Lastly, the embodiment provides an economy of scale when producing nucleic acid vaccine compositions.

Related to this embodiment, computer programs can be derived in accordance with principles in the art, which identify in a target sequence, the greatest number of cpitopes per sequence length.

5 Example 15 Polyepitopic Vaccine Compositions From Multiple Antigens

The HPV peptide epitopes of the present invention are used in conjunction with peptide epitopes from other target turnor-associated antigens to create a vaccine composition that is useful for the prevention or treatment of cancer resulting from HPV infection in multiple patients.

For example, a vaccine composition can be provided as a single polypeptide that incorporates multiple epitopes from HPV antigens as well as turnor-associated antigens that are often expressed with a target cancer, e.g., cervical cancer, associated with HPV infection, or can be administered as a composition comprising one or more discrete epitopes. Alternatively, the vaccine can be administered as a minispene construct or as dendritic cells which have been loaded with the peptide epitopes in vitro.

15 Example 16. Use of peptides to evaluate an immune response

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Peptides of the invention may be used to analyze an immune response for the presence of specific CTL or HTL populations directed to HPV. Such an analysis may be performed in a manner as that described by Ogg et al., Science 279:2103-2106, 1998. In the following example, peptides in accordance with the invention are used as a reagent for diagnostic or prognostic purposes, not as an immunogen.

In this example highly sensitive human leukocyte antigen tetrameric complexes ("tetramers") are used for a cross-sectional analysis of, for example, HPV HLA-A*0201-specific CTL frequencies from HLA A*0201-positive individuals at different stages of infection or following immunization using an HPV peptide containing an A*0201 motif. Tetrameric complexes are synthesized as described (Musey et al., N. Engl. J. Med. 337:1267, 1997). Briefly, purified HLA heavy chain (A*0201 in this example) and β2-microglobulin are synthesized by means of a prokaryotic expression system. The heavy chain is modified by deletion of the transmembrane-cytosolic tail and COOH-terminal addition of a sequence containing a BirA enzymatic biotinylation site. The heavy chain, β2-microglobulin, and peptide are refolded by dilution. The 45-kD refolded product is isolated by fast protein liquid chromatography and then biotinylated by BirA in the presence of biotin (Signa, St. Louis, Missouri), adenosine 5'triphosphate and magnesium. Streptavidin-phycocrythrin conjugate is added in a 1:4 molar ratio, and the tetrameric product is concentrated to 1 mg/ml. The resulting product is referred to as tetramer-phycocrythrin.

For the analysis of patient blood samples, approximately one million PBMCs are centrifuged at 300g for 5 minutes and resuspended in 50 µl of cold phosphate-buffered saline. Tri-color analysis is performed with the tetramer-phycocrythrin, along with anti-CD8-Tricolor, and anti-CD38. The PBMCs are incubated with tetramer and antibodies on ice for 30 to 60 min and then washed twice before formaldehyde fixation. Gates are applied to contain >>99.98% of control samples. Controls for the tetramers include both A*0201-negative individuals and A*0201-positive uninfected donors. The percentage of cells stained with the tetramer is then determined by flow cytometry. The results indicate the number of cells in the PBMC sample that contain epitope-restricted CTLs, thereby readily indicating the

extent of immune response to the HPV epitope, and thus the stage of infection with HPV, the status of exposure to HPV, or exposure to a vaccine that elicits a protective or therapeutic response.

Example 17. Use of Peptide Epitopes to Evaluate Recall Responses

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5 The peptide epitopes of the invention are used as reagents to evaluate T cell responses, such as acute or recall responses, in patients. Such an analysis may be performed on patients who have recovered from infection, who are chronically infected with HPV, or who have been vaccinated with an HPV vaccine.

For example, the class I restricted CTL response of persons who have been vaccinated may be analyzed. The vaccine may be any HPV vaccine. PBMC are collected from vaccinated individuals and HLA typed. Appropriate peptide epitopes of the invention that, optimally, bear supermotifs to provide cross-reactivity with multiple HLA supertype family members, are then used for analysis of samples derived from individuals who bear that HLA type.

PBMC from vaccinated individuals are separated on Ficoll-Histopaque density gradients (Sigma Chemical Co., St. Louis, MO), washed three times in HBSS (GIBCO Laboratories), resuspended in RPMI-1640 (GIBCO Laboratories) supplemented with L-glutamine (2nM), penicillin (50U/ml), streptomycin (50 µg/ml), and Hepes (10mM) containing 10% heat-inactivated human AB serum (complete RPMI) and plated using microculture formats. A synthetic peptide comprising an epitope of the invention is added at 10 µg/ml to each well and HBV core 128-140 epitope is added at 1 µg/ml to each well as a source of T cell help during the first week of stimulation.

In the microculture format, 4 x 10⁵ PBMC are stimulated with peptide in 8 replicate cultures in 96-well round bottom plate in 100 µl/well of complete RPMI. On day 3 and 10, 100 ul of complete RPMI and 20 U/ml final concentration of rIL-2 are added to each well. On day 7 the cultures are transferred into a 96-well flat-bottom plate and restimulated with peptide, rIL-2 and 10⁵ tradiated (3,000 rad) autologous feeder cells. The cultures are tested for cytotoxic activity on day 14. A positive CTL response requires two or more of the eight replicate cultures to display greater than 10% specific ³¹Cr release, based on comparison with uninfected control subjects as previously described (Rehermann, et al., Nature Med. 2:1104,1108, 1996; Rehermann et al., J. Clin. Invest. 97:1655-1665, 1996; and Rehermann et al., J. Clin. Invest. 98:1432-1440, 1996).

Target cell lines are autologous and allogeneic EBV-transformed B-LCL that are either purchased from the American Society for Histocompatibility and Immunogenetics (ASHI, Boston, MA) or established from the pool of patients as described (Guilhot, et al. J. Virol. 66:2670-2678, 1992).

Cytotoxicity assays are performed in the following manner. Target cells consist of either allogeneic HLA-matched or autologous EBV-transformed B lymphoblastoid cell line that are incubated overnight with the synthetic peptide epitope of the invention at 10 µM, and labeled with 100 µCi of ⁵¹Cr (Amersham Corp., Arlington Heights, IL) for 1 hour after which they are washed four times with HBSS. Cytolytic activity is determined in a standard 4-h, split well ⁵¹Cr release assay using U-bottomed 96 well plates containing 3,000 targets/well. Stimulated PBMC are tested at effect/fragret (E/T) ratios of 20-50:1 on day 14. Percent cytotoxicity is determined from the formula: 100 x ((experimental release-spontaneous release)). Maximum release is determined by

lysis of targets by detergent (2% Triton X-100; Sigma Chemical Co., St. Louis, MO). Spontaneous release is <25% of maximum release for all experiments.

The results of such an analysis indicate the extent to which HLA-restricted CTL populations have been stimulated by previous exposure to HPV or an HPV vaccine.

The class II restricted HTL responses may also be analyzed. Purified PBMC are cultured in a 96-well flat bottom plate at a density of 1.5x 10³ cells/well and are stimulated with 10 µg/ml synthetic peptide, whole antigen, or PHA. Cells are routinely plated in replicates of 4-6 wells for each condition. After seven days of culture, the medium is removed and replaced with fresh medium containing 10U/ml IL-2. Two days later, 1 µCi ³H-thymidine is added to each well and incubation is continued for an additional 18 hours. Cellular DNA is then harvested on glass fiber mats and analyzed for ³H-thymidine incorporation. Antigen-specific T cell proliferation is calculated as the ratio of ³H-thymidine incorporation in the presence of antigen divided by the ³H-thymidine incorporation in the absence of antigen.

Example 18. Induction Of Specific CTL Response In Humans

A human clinical trial for an immunogenic composition comprising CTL and HTL epitopes of the invention is set up as an IND Phase I, dose escalation study and carried out as a randomized, double-blind, placebo-controlled trial. Such a trial is designed, for example, as follows:

A total of about 27 individuals are enrolled and divided into 3 groups:

Group I: 3 subjects are injected with placebo and 6 subjects are injected with 5 µg of

20 peptide composition;

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Group II: 3 subjects are injected with placebo and 6 subjects are injected with 50 μ g peptide composition;

Group III: 3 subjects are injected with placebo and 6 subjects are injected with 500 µg of peptide composition.

25 After 4 weeks following the first injection, all subjects receive a booster inoculation at the same dosage.

The endpoints measured in this study relate to the safety and tolerability of the peptide composition as well as its immunogenicity. Cellular immune responses to the peptide composition are an index of the intrinsic activity of this the peptide composition, and can therefore be viewed as a measure of biological efficacy. The following summarize the clinical and laboratory data that relate to safety and efficacy endpoints.

Safety: The incidence of adverse events is monitored in the placebo and drug treatment group and assessed in terms of degree and reversibility.

Evaluation of Vaccine Efficacy: For evaluation of vaccine efficacy, subjects are bled

before and after nijection. Peripheral blood mononnuclear cells are isolated from fresh heparinized blood by
Ficoll-Hypaque density gradient centrifugation, aliquoted in freezing media and stored frozen. Samples are
assayed for CTL and HTL activity.

The vaccine is found to be both safe and efficacious.

Example 19. Phase II Trials In Patients Infected With HPV

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Phase II trials are performed to study the effect of administering the CTL-HTL peptide compositions to patients having cancer associated with HPV infection. The main objectives of the trials are to determine an effective dose and regimen for inducing CTLs in HPV-infected patients with cancer, to establish the safety of inducing a CTL and HTL response in these patients, and to see to what extent activation of CTLs improves the clinical picture of chronically infected HPV patients, as manifested by a reduction in viral load, e.g., the reduction and/or shrinking of lesions. Such a study is designed, for example, as follows:

The studies are performed in multiple centers. The trial design is an open-label, uncontrolled, dose escalation protocol wherein the peptide composition is administered as a single dose followed six weeks later by a single booster shot of the same dose. The dosages are 50, 500 and 5,000 micrograms per injection. Drug-associated adverse effects (severity and reversibility) are recorded.

There are three patient groupings. The first group is injected with 50 micrograms of the peptide composition and the second and third groups with 500 and 5,000 micrograms of peptide composition, respectively. The patients within each group range in age from 21-65 and represent diverse ethnic backgrounds. All of them are infected with HPV and are HIV, HCV, HBV and delta hepatitis virus (HDV) negative, but are positive for HPV DNA as monitered by PCR.

Clinical manifestations or antigen-specific T-cell responses are monitored to assess the effects of administering the peptide compositions. The vaccine composition is found to be both safe and efficacious in the treatment of HPV infection.

Example 20. Induction of CTL Responses Using a Prime Boost Protocol

A prime boost protocol similar in its underlying principle to that used to evaluate the efficacy of a DNA vaccine in transgenic mice, such as described in Example 12, can also be used for the administration of the vaccine to humans. Such a vaccine regimen can include an initial administration of, for example, naked DNA followed by a boost using recombinant virus encoding the vaccine, or recombinant protein/polypeptide or a peptide mixture administered in an adjuvant.

For example, the initial immunization may be performed using an expression vector, such as that constructed in Example 11, in the form of naked nucleic acid administered 1M (or SC or ID) in the amounts of 0.5-5 mg at multiple sites. The nucleic acid (0.1 to 1000 µg) can also be administered using a gene gun. Following an incubation period of 3-4 weeks, a booster dose is then administered. The booster can be recombinant fowlpox virus administered at a dose of 5-107 to 5x109 pld. An alternative recombinant virus, such as an MVA, canarypox, adenovirus, or adeno-associated virus, can also be used for the booster, or the polyepitopic protein or a mixture of the peptides can be administered. For evaluation of vaccine efficacy, patient blood samples will be obtained before immunization as well as at intervals following administration of the initial vaccine and booster doses of the vaccine. Peripheral blood mononuclear cells are isolated from fresh heparinized blood by Ficoll-Hypaque density gradient centrifugation, aliquoted in freezing media and stored frozen. Samples are assayed for CTL and HTL activity.

Analysis of the results indicates that a magnitude of response sufficient to achieve
40 protective immunity against HPV is generated.

Example 21. Administration of Vaccine Compositions Using Dendritic Cells (DC)

Vaccines comprising peptide epitopes of the invention can be administered using APCs, or "professional" APCs such as DC. In this example, the peptide-pulsed DC are administered to a patient to 5 stimulate a CTL response in vivo. In this method, dendritic cells are isolated, expanded, and pulsed with a vaccine comprising peptide CTL and HTL epitopes of the invention. The dendritic cells are infused back into the patient to elicit CTL and HTL responses in vivo. The induced CTL and HTL then destroy or facilitate destruction of the specific target cells that bear the proteins from which the epitopes in the vaccine are derived.

For example, a cocktail of epitope-bearing peptides is administered ex vivo to PBMC, or isolated DC therefrom. A pharmaceutical to facilitate harvesting of DC can be used, such as Progenipoietin (Monsanto, St. Louis, MO) or GM-CSF/IL-4. After pulsing the DC with peptides and prior to reinfusion into patients, the DC are washed to remove unbound peptides.

As appreciated clinically, and readily determined by one of skill based on clinical

outcomes, the number of DC reinfused into the patient can vary (see, e.g., Nature Med. 4:328, 1998; Nature
Med. 2:52, 1996 and Prostate 32:272, 1997). Although 2-50 x 10⁶ DC per patient are typically
administered, larger number of DC, such as 10⁷ or 10⁸ can also be provided. Such cell populations typically
contain between 50-90% DC.

In some embodiments, peptide-loaded PBMC are injected into patients without

Purification of the DC. For example, PBMC containing DC generated after treatment with an agent such as
Progenipoietin are injected into patients without purification of the DC. The total number of PBMC that
are administered often ranges from 10⁸ to 10¹⁰. Generally, the cell doses injected into patients is based on
the percentage of DC in the blood of each patient, as determined, for example, by immunofluorescence
analysis with specific anti-DC antibodies. Thus, for example, if Progenipoietin™ mobilizes 2% DC in the
peripheral blood of a given patient, and that patient is to receive 5 x 10⁶ DC, then the patient will be injected
with a total of 2.5 x 10⁸ peptide-loaded PBMC. The percent DC mobilized by an agent such as
Progenipoieti™ is typically estimated to be between 2-10%, but can vary as appreciated by one of skill in
the art

30 Ex vivo activation of CTL/HTL responses

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Alternatively, ex vivo CTL or HTL responses to HPV antigens can be induced by incubating in tissue culture the patient's, or genetically compatible, CTL or HTL precursor cells together with a source of APC, such as DC, and the appropriate immunogenic peptides. After an appropriate incubation time (typically about 7-28 days), in which the precursor cells are activated and expanded into effector cells, the cells are infused back into the patient, where they will destroy (CTL) or facilitate destruction (HTL) of their specific target cells, t.e., tumor cells.

Example 22. Alternative Method of Identifying Motif-Bearing Peptides

Another method of identifying motif-bearing peptides is to elute them from cells bearing

40 defined MHC molecules. For example, EBV transformed B cell lines used for tissue typing have been

extensively characterized to determine which HLA molecules they express. In certain cases these cells express only a single type of HLA molecule. These cells can be infected with a pathogenic organism or transfected with nucleic acids that express the antigen of interest, eg. HPV regulatory or structural proteins. Peptides produced by endogenous antigen processing of peptides produced consequent to infection (or as a result of transfection) will then bind to HLA molecules within the cell and be transported and displayed on the cell surface. Peptides are then eluted from the HLA molecules by exposure to mild acid conditions and their amino acid sequence determined, e.g., by mass spectral analysis (e.g., Kubo et al., J. Immunol. 152:3913, 1994). Because the majority of peptides that bind a particular HLA molecule are motif-bearing, this is an alternative modality for obtaining the motif-bearing peptides correlated with the particular HLA molecule expressed on the cell.

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Alternatively, cell lines that do not express endogenous HLA molecules can be transfected with an expression construct encoding a single HLA allele. These cells can then be used as described, i.e., they can be infected with a pathogen or transfected with nucleic acid encoding an antigen of interest to isolate peptides corresponding to the pathogen or antigen of interest that have been presented on the cell surface. Peptides obtained from such an analysis will bear motif(s) that correspond to binding to the single HLA allele that is expressed in the cell.

As appreciated by one in the art, one can perform a similar analysis on a cell bearing more than one HLA allele and subsequently determine peptides specific for each HLA allele expressed. Moreover, one of skill would also recognize that means other than infection or transfection, such as loading with a protein antigen, can be used to provide a source of antigen to the cell.

The above examples are provided to illustrate the invention but not to limit its scope. For example, the human terminology for the Major Histocompatibility Complex, namely HLA, is used throughout this document. It is to be appreciated that these principles can be extended to other species as well. Thus, other variants of the invention will be readily apparent to one of ordinary skill in the art and are encompassed by the appended claims. All publications, patents, and patent application cited herein are hereby incorporated by reference for all purposes.

TABLE I

SUPERMOTIFS	POSITION	POSITION	POSITION
	2 (Primary Anchor)	3 (Primary Anchor)	C Terminus (Primary
	_ (,	`	Anchor)
A1	T, I, L, V, M, S		F, W, Y
A2	L, I, V, M, A, T, Q		I, V, M, A, T, L
A3	V, S, M, A, T, L, I		R,K
A24	Y, F, W, I, V, L, M, T		F, I, Y, W, L, M
B7	P		V, I, L, F, M, W, Y, A
B27	R, H, K		F, Y, L, W, M, I, V, A
B44	E , <i>D</i>		F, W, L, I, M, V, A
B58	A, T, S		F, W, Y, L, I, V, M, A
B62	Q, L, I, V, M, P		$\mathbf{F}, \mathbf{W}, \mathbf{Y}, M, I, V, L, A$
MOTIFS			
A1	T, S, M		Y
A1		D , E , <i>A</i> , <i>S</i>	Y
A2.1	L, M, V, Q, I, A, T		V, L, I, M, A, T
A3	L, M, V, I, S, A, T, F,		K, Y, R, H, F, A
	C, G, D		
A11	V, T, M, L, I, S, A,		K, R, Y, H
	G , N , <i>C</i> , <i>D</i> , <i>F</i>		
A24	Y, F, W, M		F, L, I, W
A*3101	M, V, T, A, L, I, S		R, K
A*3301	M, V, A, L, F, I, S, T		R, K
A*6801	A, V, T, M, S, L, I		R, K
B*0702	P		L, M, F, W, Y, A, I, V
B*3501	P		L, M, F, W, Y, I, V, A
B51	P		L, I, V, F, W, Y, A, M
B*5301	P		I, M, F, W, Y, A, L, V
B*5401	P		\mathbf{A} , \mathbf{T} , \mathbf{I} , \mathbf{V} , L , M , F ,
			W, Y

Bolded residues are preferred, italicized residues are less preferred: A peptide is considered motif-bearing if it has primary anchors at each primary anchor position for a motif or supermotif as specified in the above table.

TABLE Ia

SUPERMOTIFS	POSITION	POSITION	POSITION
	2 (Primary Anchor)	3 (Primary Anchor)	C Terminus (Primary
			Anchor)
Al	T, I, L, V, M, S		F, W, Y
A2	V, Q, A, T		I, V, L, M, A, T
A3	V, S, M, A, T, L, I		R, K
A24	Y, F, W, I, V, L, M, T		F, I, Y, W, L, M
B7	P		V, I, L, F, M, W, Y, A
B27	R, H, K		F, Y, L, W, M, I, V, A
B58	A, T, S		F, W, Y, L, I, V, M, A
B62	Q, L, I, V, M, P		F, W, Y, M, I, V, L, A
MOTIFS			
A1	T, S, M		Y
A1		D , E,A, S	Y
A2.1	V, Q, A, T*		V, L, I, M, A, T
A3.2	L, M, V, I, S, A, T, F,		K, Y, R, H, F, A
	C, G, D		
A11	V, T, M, L, I, S, A,		K, R, H, Y
1	G, N, C, D, F		
A24	Y ,F, W		F, L, I, W

^{*}If 2 is V, or O, the C-term is not L

Bolded residues are preferred, italicized residues are less preferred: A peptide is considered motif-bearing if it has primary anchors at each primary anchor position for a motif or supermotif as specified in the above table.

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TABLEII

						NOILLISON	NO			
			Z	<u> </u>	4	3	9	4	SSI	C-terminus
SUPE	SUPERMOTIFS									
A L			1° Anchor T,I,L,V,M,S							1º Anchor F,W,Y
A2			L,I,V,M,A,			•				L,I,V,M,A,T
£4	preferred		1° Anchor V,S,M,A,T, L,I	Y,F,W, (4/5)		-	Y,F,W, (3/5)	Y,F,W, (4/5) P, (4/5)	P, (4/5)	P,K
	deleterious	D,E (3/5); P, (5/5)		D,E, (4/5)						
A24			1° Anchor Y,F,W,I,V,							1° Anchor F,1,Y,W,L,M
B7	preferred	F,W,Y (5/5) L,I,V,M, (3/5)	1°Anchor P	F,W,Y (4/5)					F,W,Y, (3/5)	1°Anchor V,I,L,F,M.W.Y.A
	deleterious	D,E (3/5); P(5/5); G(4/5); A(3/5); Q,N, (3/5)				D,E, (3/5)) G, (4/5)	Q,N, (4/5)	D,E, (4/5)	
B27			1° Anchor R.H.K					., .,		I. Anchor F,Y,L,W,M,V,A
B44			1º Anchor E,D							I. Anchor F,W,Y,L,I,M,V,A
B58			1º Anchor A,T,S							I. Anchor F,W,Y,L,I,V,M,A
B62			1° Anchor Q.L.J.V.M. P						•	1° Anchoi F, W, Y, M, I, V, L, A
				-						

	•					POSITION		×		
		1	Ø	<u></u>	40	<u> </u>	9	0	20	C-terminus
MOTIFS	ES									
A l 9-mer	preferred	G,F,Y,W,	1°Anchor S,T,M,	D,E,A,	Y,F,W,		ď.	D,E,Q,N, Y,F,W,	Y,F,W,	1°Anchor Y
	deleterious D,E,	a'd		R,H,K,L,1,V A, M,P,	₹	ΰ	¥			
A 1 9-mer	ргегетед	G,R,H,K	A,S,T,C,L,I V,M,	A,S,T,C,L,I <u>l'Anchor</u> V,M, D,E,A,S	G,S,T,C,		A,S,T,C,	A,S,T,C, L,I,V,M,	D,E,	1°Anchor Y
	deleterious A	< .	R,H,K,D,E, P,Y,F,W,		D,E,	P,Q,N,	R,H,K,	P,G,	G,P,	

						POSITION	z				
			Z	<u> </u>	4	<u> </u>	Ø			ত ক ব ক্র	C-terminus
A1 10-mer	peferred	Y,F,W,	1°Anchor S.T.M	D,E,A,Q,N,	. ⊀	Y,F,W,Q,N,		P,A,S,T,C, G,D,E,	G,D,E,	C-terminus P,	1°Anchor Y
	deleterious	G,P,		R,H,K,G,L,I D,E, V,M,	D,E,	к,н,к,	O,N,O	R,H,K,Y,F, R,H,K, W,	R,H,K,	<	
A1 10-mer	preferred	Y,F,W,	S,T,C,L,I,V M,	1°Anchor D,E,A,S	ŕ	Y,F,W,		P,G,	σ̂	Y,F,W,	l°Anchor Y
	deleterious	R,H,K,	R,H,K,D,E, P,Y,F,W,			of .	ď		P,R,H,K, Q,N,	z'. Ö	
A2.1 9-mer	preferted	Y,F,W,	1° Auchor L,M,I.P.Q.	Y,F,W,	s,r,c,	Y,F,W,		₹	<u>-</u>	1°Anchar V,L,I,M,A,T	
	deleterious	D,E,P,		р, Е, К, Н			к,к,н	D,E,R,K,H			
A2.1 10-me	A2.1 preferred 10-mer	A,Y,F,W,	L,M,J,V,Q,	L,V,I,M,	σ̈́		Ů.		F,Y,W,L, V,I,M,		1°Anchor V,L,I,M,A,T
	deleterious	deleterious D.E.P.,		D,E,	R,K,H,A,	p.°		R,K,H,	D,E,R,K, R,K,H, H,	R,K,H,	

	1										
			Ø	6	(4)	[2]	S		(20)	<u>ල</u> ා ප	C- terminus
	preferred	R,H,K,	1°Anchor L,M,V,I,S, A,T,F,C,G D	Y,F,W,	P,R,H,K,Y, F,W,	₹	Y,F,W,		٠. د	C-terminus 1ºAnchor K, Y,R, H, F, A	
	deleterious	D,E,P,		D,E							
I V	preferred	ď.	l*Anchor V,T,L,M,I, S,A,G,N,C, D,F	Y,F,W,	Y,FW,	. 4	Y,F,W,	Y,FW,	e,	L*Anchor K,,RY,H	
	deleterious	D,E,P,						<	ú		
A24 9-mer	preferred	Y,F,W,R,H,K,	1°Anchor Y,F,W,M		S,T,C			Y,F,W,	Y,F,W,	1°Anchor F,L,1,W	
	deleterious D,E,G,	D,E,G,		, E, C	oʻ	Q,N,P,	D,E,R,H,K,	ග්	A,Q,N,		
A24 10-mer	preferred		1°Anchor Y,F,W,M		o,	Y,F,W,P,		ď.			I Anchor F,L,I,W
	deleterious			G,D,E	N,Q	R,H,K	D,E	. 4	O,N,	D,E,A,	
101	A3101 preferred	R,H,K,	1°Anchor M,V,T,A.L, I,S	Y,F,W,	ď.		Y,F,W,	Y,F,W,	A,P,	1°Anchor R,K	
	deleterious	D,E,P,		D,E,		A,D,E,	D,E,	D,E,	Ξ'Q		

ප					,r ,Y,A,		7'A')	
<u>a</u>	or C-terminus 1ºAnchor R,K		1°Anchor R,K		L'Anchor L,M,F,W,Y,A, I,V		!*Anchor L,M,F,W,Y,,, F,A	
<u></u>	D		a.°	ď.	P,A	· D,E,		
E	A,Y,F,W		Y,F,W,		R,H,K,	zî Ø	F,W,Y,	
<u> </u>					R,H,K,	G,D,E,		Ġ
2			Y,F,W,L,I, V,M	R,H,K,	R,H,K,	D,E,		Ö
40	ı					D,E,		
<u>e</u>		D,E		D,E,G,	R,H,К ,	D,E,P,	F,W,Y,	
(ZI	Anchor ,V,A,L,F,		1°Anchor A,V,T,M,S, L,I		L*Anchor		1°Anchor P	
		g,P	Y,F,W,S,T,C,	G,P,	R,H,K,F,W,Y,	D,E,Q,N,P,	F,W,Y,L,I,V,M,	A,G,P,
	A3301 preferred	deleterious G,P	A6801 preferred	deleterious	B0702 preferred	deleterious	B3501 preferred	deleterious A,G,P,
	A3301		A6801		B0702		B3501	

						POSTTION	_				
		1	Z	කෙ	(2 2)	S	<u> </u>		.	ි කො ලකු පි	C- terminus
BSį	ргебетед	L,I,V,M,F,W,Y,	1°Anchor P	F,W,Y,	S,T,C,	F,W,Y,		ď	F,W,Y,	L'Anchor L'I,V,F,W,	
	deleterious	deleterious A,G,P,D,E,R,H,K, S,T,C,				D,E,	Ġ	D,E,Q,N,	G,D,E,		
B5301	B5301 preferred	L,,,v,,d,F,W,Y,	L*Anchor P	F,W,Y,	s,T,C,	F,W,Y,		L,I,V,M,F, W,Y,	F,W,Y,	1°Anchor 1,M,F,W,Y,	
	deleterious	A,G,P,Q,N,				-	ů	R,H,K,Q,N, D,E,	D,E,		
B5401	B5401 preferred	F,W,Y,	1°Anchor P	F,W,Y,L,I,V M,		L,I,V,M,	•	A,L,I,V,M,	F,W,Y,A,]	A,L,I,V,M, F,W,Y,A,P, L*Anchor A,I'I,I,V,L, M,F,W,Y	
	deleterious	deleterious G.P.Q.N.D.E,		G,D,E,S,T,C,		R,H,K,D,E, D,E,	D,E,	Q,N,D,G,E,	D,E,	.	

Italicized residues indicate less preferred or "obtrated" residues.
The information in Table II is specific for 9-mers unless otherwise specified.
Secondary anchor specificities are designated for each position independently.

Table III	Е					POSITION				
MOTIFS	S)	1° anchor I	図	<u>B</u>	3	2	1° anchor 6	<u> </u>	2	<u> </u>
DR4	preferred	F, M, Y, L, I, V, W,	Σ	T,		1	V, S, T, C, P, A, I, I, M,	М, Н,		М, Н
	deleterious				w,			સ		W, D, E
DR1	preferred	M, F, L, I, V, W, Y,			P, A, M, Q,		V, M, A, T, S, P, M, L, I, C,	Ř		A, V, M
	deleterious		o	С, н	F, D	C, W, D		G, D, E,	Q	
DR7	preferred	M, F, L, I, V, W, Y,	Μ,	»́	ď.		I, V, M, S, A, C, T, P, L,	Ř		I, V
	deleterious		౮		ී			G, R, D,	z	O
R S.	DR Supermotif	M, F, L, I, V, W, Y,					V, M, S, T, A, C, P, L, I,			
83	DR3 MOTIFS	1° anchor 1	Z	<u>rea</u>	1° anchor 4	S	l° anchor 6			
motif a preferred	Pa	L, I, V, M, F, Y,			Q					
motif b preferred	, pp	L, I, V, M, F, A, Y,			D, N, Q, E, S, T		К, R, Н			

Italicized residues indicate less preferred or "tolerated" residues. Secondary anchor specificities are designated for each position independently.

Table IV: HLA Class I Standard Peptide Binding Affinity.

ALLELE	STANDARD	SEQUENCE	SEQ ID	STANDARD
	PEPTIDE		NO:	BINDING
				AFFINITY (nM)
A*0101	944.02	YLEPAIAKY	51487	25
A*0201	941.01	FLPSDYFPSV	51488	5.0
A*0202	941.01	FLPSDYFPSV	51488	4.3
A*0203	941.01	FLPSDYFPSV	51488	10
A*0205	941.01	FLPSDYFPSV	51488	4.3
A*0206	941.01	FLPSDYFPSV	51488	3.7
A*0207	941.01	FLPSDYFPSV	51488	23
A*6802	1072.34	YVIKVSARV	51489	8.0
A*0301	941.12	KVFPYALINK	51490 .	11
A*1101	940.06	AVDLYHFLK	51491	6.0
A*3101	941.12	KVFPYALINK	51490	18
A*3301	1083.02	STLPETYVVRR	51492	29
A*6801	941.12	KVFPYALINK	51490	8.0
A*2402	979.02	AYIDNYNKF	51493	12
B*0702	1075.23	APRTLVYLL	51494	5.5
B*3501	1021.05	FPFKYAAAF	51495	7.2
B51	1021.05	FPFKYAAAF	51495	5.5
B*5301	1021.05	FPFKYAAAF	51495	9.3
B*5401	1021.05	FPFKYAAAF	51495	10

Table V. HLA Class II Standard Peptide Binding Affinity.

Allele	Nomenclature	Standard	Sequence	SEQ ID	Binding
•		Peptide		NO:	Affinity
					(nM)
DRB1*0101	DR1	515.01	PKYVKQNTLKLAT	51496	5.0
DRB1*0301	DR3	829.02	YKTIAFDEEARR	51497	300
DRB1*0401	DR4w4	515.01	PKYVKQNTLKLAT	51496	45
DRB1*0404	DR4w14	717.01	YARFQSQTTLKQKT	51498	. 50
DRB1*0405	DR4w15	717.01	YARFQSQTTLKQKT	51498	38
DRB1*0701	DR7	553.01	QYIKANSKFIGITE	51499	25
DRB1*0802	DR8w2	553.01	QYIKANSKFIGITE	51499	49
DRB1*0803	DR8w3	553.01	QYIKANSKFIGITE	51499	1600
DRB1*0901	DR9	553.01	QYIKANSKFIGITE	51499	75
DRB1*1101	DR5w11	553.01	QYIKANSKFIGITE	51499	20
DRB1*1201	DR5w12	1200.05	EALIHQLKINPYVLS	51500	298
DRB1*1302	DR6w19	650.22	QYIKANAKFIGITE	51499	3.5
DRB1*1501	DR2w2β1	507.02	GRTQDENPVVHFFKNIV	51501	9.1
			TPRTPPP		
DRB3*0101	DR52a	511	NGQIGNDPNRDIL	51502	470
DRB4*0101	DRw53	717.01	YARFQSQTTLKQKT	51498	58
DRB5*0101	DR2w2β2	553.01	QYIKANSKFIGITE	51499	20

Table VI

	Allelle-specific HLA-supertype members	ype members
HLA-supertype	Verified*	Predicted
IV	A*0101, A*2501, A*2601, A*2602, A*3201	A*0102, A*2604, A*3601, A*4301, A*8001
A2	A*0201, A*0202, A*0203, A*0204, A*0205, A*0206, A*0207, A*0209, A*0214, A*6802, A*6901	A*0208, A*0210, A*0211, A*0212, A*0213
A3	A*0301, A*1101, A*3101, A*3301, A*6801	A*0302, A*1102, A*2603, A*3302, A*3303, A*3401, A*3402, A*6601, A*6602, A*7401
A24	A*2301, A*2402, A*3001	A*2403, A*2404, A*3002, A*3003
B7	B*0702, B*0703, B*0704, B*0705, B*1508, B*3501, B*3502, B*3503, B*3503, B*3501, B*3502, B*3501, B*3503, B*3503, B*3503, B*3503, B*5103, B*5105, B*5105, B*5105, B*55103, B*551	B*1511, B*4201, B*5901
B27	B*1401, B*1402, B*2702, B*2703, B*2704, B*2705, B*2706, B*3801, B*3901, B*3902, B*7301	B*2701, B*2707, B*2708, B*3802, B*3903, B*3904, B*3905, B*4801, B*4802, B*1510, B*1518, B*1503
B44	B*1801, B*1802, B*3701, B*4402, B*4403, B*4404, B*4001, B*4002, B*4006	B*4101, B*4501, B*4701, B*4901, B*5001
B58	B*5701, B*5702, B*5801, B*5802, B*1516, B*1517	
B62	B*1501, B*1502, B*1513, B*5201	B*1301, B*1302, B*1504, B*1505, B*1506, B*1507,

Verified alleles include alleles whose specificity has been determined by pool sequencing analysis, peptide binding assays, or by analysis of the sequences of CTL epitopes. æ

B*1301, B*1302, B*1504, B*1505, B*1506, B*1507, B*1515, B*1520, B*1521, B*1512, B*1514, B*1510

Predicted alleles are alleles whose specificity is predicted on the basis of B and F pocket structure to overlap with the supertype specificity. 6

Table VII HLA-A1 Supermotif Peptides

1	2	3	4	HPV16	E1	8	441
HPV16	E1	10	206	HPV16	E1	10	419
HPV16	E1	8	524	HPV16	E1	10	118
HPV16	E1	9	82	HPV16	E1	8	343
HPV16	E1	11	353	HPV16	E1	10	125
HPV16	E1	10	368	HPV16	E1	11	582
HPV16	E1	11	41	HPV16	E1	8	313
HPV16	El	8	372	HPV16	E1	9	313
HPV16	E1	10	249	HPV16	E1	10	313
HPV16	E1	9	43	HPV16	E1	8	432
HPV16		9	384	HPV16	E1	9	250
HPV16	E1	10	603	HPV16	E1	10	484
HPV16	E1	11	603	 HPV16	E1	8	421
HPV16	E1	8	356	HPV16 HPV16	E1 E1	9	314 314
HPV16 HPV16	E1 E1	10 9	356 63	HPV16	E1	11	231
HPV16	E1	9	152	HPV16	E1	9	253
HPV16	E1	9	331	HPV16	E1	11	498
HPV16	E1	8	51	HPV16	E1	11	345
HPV16		9	493	HPV16	E1	11	443
HPV16	E1	9	445	HPV16	E1	10	217
HPV16	E1	8	456	HPV16	E1	9	584
HPV16	El	11	453	HPV16	E1	11	584
HPV16	E1	8	219	HPV16	E1	8	274
HPV16	El	9	586	HPV16	E1	11	261
HPV16	E1	8	501	HPV16	E1	9	578
HPV16	E1	9	501	HPV16	E1	11	578
HPV16	E1	11	466	HPV16	E2	11	331
HPV16	E1	9	325	HPV16	E2	11	41
HPV16	E1	11	519	HPV16	E2	8	314
HPV16	E1	10	272	HPV16	E2	11	309
HPV16	E1	9	163	HPV16	E2	8	124
HPV16	E1	8	571	HPV16	E2	11	124
HPV16		8	12	HPV16	E2	8	25
HPV16		9	12	HPV16	E2	9 .	25
HPV16	E1	11	216	HPV16	E2	9	263
HPV16	E1	9	263	HPV16	E2	9	338
HPV16	E1	8	348	HPV16 HPV16	E2 E2	11	22 74
HPV16	E1	11	329	HPV16	E2	8	80
HPV16	E1 E1	9	326 369	HPV16	E2	11	168
HPV16 HPV16	E1	11	369	HPV16	E2	9	163
HPV16	E1	10	311	HPV16	E2	9	35
HPV16		11	311	HPV16	E2	10	35
HPV16	E1	8	610	HPV16	E2	8	193
HPV16	El	11	483	HPV16	E2	10	332
HPV16	E1	10	227	HPV16	E2	9	329
HPV16	E1	11	323	HPV16	E2	9	354
HPV16	E1	10	252	HPV16	E2	11	77
HPV16	E1	8	254	HPV16	E2	9	84
HPV16	E1	9	357	HPV16	E2	8	296
HPV16	E1	11	48	HPV16	E2	10	296
HPV16	E1	10	583	HPV16	E2	8	127
HPV16	E1	9	207	HPV16	E2	11	9
HPV16	E1	10	520	HPV16	E2	10	106
HPV16	E1	10	454	HPV16	E2	8	76
HPV16	E1	9	420	HPV16	E2	8	151
HPV16	E1	9	273	HPV16	E2	9	151
HPV16	Eļ	10	567	HPV16	E2	10	191
HPV16	E1	10	600	HPV16	E2	8	37

Table VII HLA-A1 Supermotif Peptides

HPV16	E2	10	23	HPV16	E6	10	79
HPV16	E2	11	23	HPV16	E6	9	44
HPV16	E2	11	261	HPV16	E6	11	44
HPV16	E2	11	144	HPV16	E6	8	4.3
HPV16	E2	8	355	HPV16	E6	10	43
HPV16	E2	10	78	HPV16	E6	11	89
HPV16		9	297	HPV16	E6	11	29
HPV16		10	93	HPV16	E6	10	77
HPV16		8	334	HPV16	E7	10	14
HPV16		10	310	HPV16	E7	8	4
HPV16		11	128	HPV16		8	18
HPV16		9	146	HPV16		9	373
HPV16		10	146	HPV16		11	292
HPV16		9	192	HPV16		10	251
HPV16		9	333	HPV16		9	249
HPV16		10	145	HPV16		11	484
HPV16		11	145	HPV16		8	154
HPV16		8	147	HPV16		9	228
HPV16		9	147	HPV16		8	17
HPV16		11	92	HPV16		9	17
HPV16		8	312	HPV16		9	378
HPV16		10	312	HPV16		8	474
HPV16		8	131	HPV16		10	5
HPV16		9	159	HPV16		8	481
HPV16		10	159	HPV16		9	348
HPV16		10	54	HPV16		8	499
HPV16		9	7	HPV16		11	323
	E5	11	5	HPV16		11	307
HPV16		9	60	HPV16		9	438
HPV16		9	72	HPV16		9	22
HPV16		9	64	HPV16		8	102
HPV16		8	43	HPV16		10	102
	E5	10	51	HPV16		11	418
HPV16		8	61	HPV16		11	86
HPV16		8	73	HPV16		8	374
HPV16		9	42	HPV16		11	11
HPV16	E5	9	11	HPV16		10	407
HPV16		8	32	HPV16		11	406
HPV16		11	47	HPV16		11	151
HPV16		10	48	HPV16		10	90
HPV16		11	70	HPV16		8	46
HPV16		9	31			8	68
HPV16		10	41	HPV16		9	68
HPV16		8	8	HPV16		8	409
HPV16		10	10	HPV16		10	87
	E5	11	40	HPV16			226
HPV16	E5	11	9				263
HPV16		8	50	HPV16		9	325
HPV16		11	50	HPV16		8	311
HPV16		10	63			8	421
HPV16		9	68			10	421
HPV16		10	68	HPV16		11	247
HPV16		10	58	HPV16			466
		11	73	HPV16			43
HPV16			32	HPV16			331
		8	32 92	HPV16			280
HPV16			92 125	HPV16		10	100
HPV16		9	125 80	HPV16		9	67
HPV16 HPV16		9	80 59			9 10	67
		8	79	HPV16			253
HPV16	E0	0	19	ULATO	H.L		4.13

Table VII HLA-A1 Supermotif Peptides

			 -p		
HPV16 L1	11	28	HPV16 L2	9	429
HPV16 L1	10	419	HPV16 L2	10	124
HPV16 L1	10	324	HPV16 L2	8	386
HPV16 L1	10	308	HPV16 L2	11	383
HPV16 L1	11	308	HPV16 L2	10	172
HPV16 L1	9.	422	HPV16 L2	9	358
HPV16 L1	8	423	HPV16 L2	8	221
HPV16 L1	8	439	HPV16 L2	11	44
HPV16 L1	9	408	HPV16 L2	8	342
HPV16 L1	11	327	HPV16 L2	9	234
HPV16 L1	11	376	HPV16 L2	11	9
HPV16 L1	9	252	HPV16 L2	8	319
HPV16 L1	11	65	HPV16 L2	9	319
HPV16 L1	8	379	HPV16 L2	10	319
HPV16 L1	11	379	HPV16 L2	10	274
HPV16 L1	10	264	HPV16 L2 HPV16 L2	10 9	360 125
HPV16 L1	11	264	HPV16 L2	11	104
HPV16 L1 HPV16 L1	9 10	91 44	HPV16 L2	8	107
HPV16 L1	8	326	HPV16 L2	10	184
HPV16 L1	9	30	HPV16 L2	9	185
HPV16 L1	9	260	HPV16 L2	8	186
HPV16 L1	8	7	HPV16 L2	10	384
HPV16 L1	8	389	HPV16 L2	9	40
HPV16 L1	8	275	HPV16 L2	9	438
HPV16 L1	8	53	HPV16 L2	10	438
HPV16 L1	9	53	HPV16 L2	8	399
HPV16 L2	11	356	HPV16 L2	8	359
HPV16 L2	11	293	HPV16 L2	11	359
HPV16 L2	8	261	HPV16 L2	9	295
HPV16 L2	10	340	HPV16 L2	8	156
HPV16 L2	11	242	HPV16 L2	9	398
HPV16 L2	9	259	HPV16 L2	9	244
HPV16 L2	10	259	HPV16 L2	11	153
HPV16 L2	10	364	HPV16 L2	10	154
HPV16 L2	10	63	HPV16 L2	9	106 155
HPV16 L2	11	218	HPV16 L2 HPV16 L2	9 10	393
HPV16 L2	8	65	HPV16 L2	10	437
HPV16 L2 HPV16 L2	8	439 439	HPV16 L2	11	437
HPV16 L2	10	45	HPV18 E1	10	213
HPV16 L2	11	45	HPV18 E1	11	526
HPV16 L2	10	243	HPV18 E1	11	40
HPV16 L2	8	250	HPV18 E1	8 .	531
HPV16 L2	8	430	HPV18 E1	9	531
HPV16 L2	10	105	HPV18 E1	11	216
HPV16 L2	10	248	HPV18 E1	10	437
HPV16 L2	9	318	HPV18 E1	9	240
HPV16 L2	10	318	HPV18 E1	8	363
HPV16 L2	11	318	HPV18 E1	10	363
HPV16 L2	10	39	HPV18 E1	9	391
HPV16 L2	8	323	HPV18 E1	10	637
HPV16 L2	11	427	HPV18 E1	9	42
HPV16 L2	9	249	HPV18 E1	10	610
HPV16 L2	11	183	HPV18 E1	11	610 62
HPV16 L2	10	294	HPV18 E1 HPV18 E1	9 10	62 375
HPV16 L2	11	454	HPV18 E1	8	375
HPV16 L2	8	276 273	HPV18 E1	9	587
HPV16 L2	11	397	HPV18 E1	9	338
HPV16 L2	±0	33/	WIATO DI	-	550

Table VII HLA-A1 Supermotif Peptides

HPV18	E1	8	50	HPV18	E1	9	591
HPV18	E1	9	500	HPV18		11	591
HPV18	E1	11	460	HPV18	E1	11	505
HPV18	E1	8	463	HPV18	E1	9	81
HPV18	E1	10	399	HPV18	E1	9	280
HPV18	E1	9	452	HPV18		8	339
HPV18	E1	8	226	HPV18	E1	9	585
HPV18	E1	8	130	HPV18	E1	11	585
HPV18	E1	8	508	HPV18	E2	10	82
HPV18	E1	9	508	HPV18	E2	10	154
HPV18	E1	11	223	HPV18	E2	11	154
HPV18	E1	8	11	HPV18	E2 E2	11	132
	E1	9	11	HPV18 HPV18	E2 E2	10 8	14 156
HPV18 HPV18	E1 E1	10 10	473 279	HPV18	E2	9	156
HPV18		10	249	HPV18	E2	8	29
HPV18	E1	9	270	HPV18	E2	9	29
HPV18	E1	11	352	HPV18	E2	8	315
HPV18	E1	11	336	HPV18	E2	11	26
HPV18	E1	10	506	HPV18	E2	9	354
HPV18	E1	11	506	HPV18	E2	11	104
HPV18	E1	10	461	HPV18	E2	9	161
HPV18	E1	10	590	HPV18	E2	9	338
HPV18	E1	8	439	HPV18	E2	9	329
HPV18	E1	10	318	HPV18	E2	9	39
HPV18	E1	11	318	HPV18	E2	10	39
HPV18	E1	10	234	HPV18	E2	10	133
HPV18	E1	8	401	HPV18	E2	11	133
HPV18	E1	8	490	HPV18	E2	8	297
HPV18	E1	11	490	HPV18	E2	8	107
HPV18	E1	10	259	HPV18	E2	9	185
HPV18	E1	8	281	HPV18 HPV18	E2 E2	10	33 38
HPV18	E1	8	261	HPV18	E2	10 11	38
HPV18	E1 E1	9	364 224	HPV18	E2	9	220
HPV18 HPV18	E1	10 9	376	HPV18	E2	9	88
HPV18	E1	11	376	HPV18	E2	11	56
HPV18	E1	9	214	HPV18	E2	9	305
HPV18	E1	10	527	HPV18		11	230
	E1	11	47	HPV18	E2	8	233
	E1	10	574	HPV18	E2	8	355
	E1	8	428	HPV18	E2	11	140
HPV18	E1	11	487	HPV18	E2	10	57
HPV18	E1	8	448	HPV18	E2	10	97
	E1	10	607	HPV18		10	231
	E1	10	426	HPV18	E2	8	157
HPV18	E1	10	80	HPV18	E2	9	232
	E1	11	589	HPV18	E2	11	96
	E1	10	128	HPV18	E2	11	173
	E1	8	320	HPV18 HPV18	E2 E2	8	143
	E1	9	320	HPV18	E2	9	135
HPV18 HPV18	E1 E1	10	320 321	HPV18	E2	9	164
	E1	9	321	HPV18	E2	10	164
	E1	8	321	HPV18	E5	9	47
	E1	9	260	HPV18	E5	11	47
HPV18	E1	11	238	HPV18	E5	11	27
	E1	10	533	HPV18	E5	10	6
	E1	8	532	HPV18	E5	8	50
	E1	11	532	HPV18	E5	8	43

Table VII HLA-A I Supermotif Peptides

				•	•			
HPV18	E5	11	43		HPV18	L1	9	383
HPV18	E5	11	40		HPV18	L1	9	175
HPV18	E5	10	22		HPV18	L1	10	175
HPV18	E5	9	2		HPV18		8	38
HPV18		8	1		HPV18		10	13
HPV18		10	1		HPV18		11	454
HPV18		11	21		HPV18		9	428
HPV18		8	24		HPV18		11	428
HPV18		10	24		HPV18		10	40
HPV18		8	3		HPV18 HPV18		11 11	39 46
HPV18		9	25 44		HPV18		10	47
HPV18 HPV18		10 9	4.2		HPV18		10	9
HPV18		10	41		HPV18		10	443
HPV18		8	27		HPV18		9	360
HPV18		10	77		HPV18		10	125
HPV18		8	40		HPV18		11	8
HPV18		10	40		HPV18	L1	9	14
HPV18	E6	11	43		HPV18	L1	8	103
HPV18	E6	8	120		HPV18	L1	9	103
HPV18	E6	11	117		HPV18		8	445
HPV18	E6	8	92		HPV18		8	104
HPV18	E6	10	36		HPV18		11	298
HPV18		9	41		HPV18		11	261
HPV18		8	74		HPV18		10	36
HPV18		11	24		HPV18		8	457
HPV18		11	89		HPV18		10	457 510
HPV18		9 11	37 37	-	HPV18		8	52
HPV18 HPV18		8	38		HPV18		9	57
HPV18		10	38		HPV18		11	282
HPV18		10	72		HPV18		11	173
HPV18		9	82		HPV18	L1	8	28
HPV18		10	77		HPV18	L1	10	26
HPV18	E7	11	90		HPV18	L1	9	472
HPV18	E7	9	92		HPV18		11	472
HPV18	E7	9	88		HPV18		11	412
HPV18		9	78		HPV18		8	315
HPV18		8	93		HPV18		8	366
HPV18		11	63		HPV18		9	137 287
HPV18		8	345		HPV18 HPV18		8	410
HPV18 HPV18		11 8	407 310		HPV18		9	102
HPV18		11	2		HPV18		10	102
HPV18		9	284		HPV18		10	135
HPV18		8	122		HPV18		8	81
HPV18		10	122		HPV18	L1	8	288
HPV18		11	520		HPV18	L1	8	459
HPV18		9	364		HPV18	L1	10	359
HPV18	L1	10	364		HPV18		8	475
HPV18	L1	9	263		HPV18		10	455
HPV18		8	330		HPV18		9	458
HPV18		10	203		HPV18		11	100
HPV18		8	49		HPV18		10	408
HPV18		11	49		HPV18		11	78 442
HPV18		8	517		HPV18		11 9	444
HPV18 HPV18		8	145 177		HPV18		11	327
HPV18		11	342		HPV18		11	362
HPV18		11	358		HPV18		9	474
77E A T O		**	550			-	-	

Table VII HLA-A1 Supermotif Peptides

HPV18	L1	8	473	HPV18 L2	11	364
HPV18	Ll	10	473	HPV18 L2	8	220
HPV18	L1	9	126	HPV18 L2	10	450
HPV18	L1	8	89	HPV18 L2	10	247
HPV18	L1	8	361	HPV18 L2	11	246
HPV18	L1	9	295	HPV18 L2	8	393
HPV18	L1	11	35	HPV18 L2	11	147
HPV18	L1	8	425	HPV18 L2	10	153
HPV18		9	4	HPV18 L2	8	365
HPV18	L1	8	88	HPV18 L2	10	365
HPV18		9	88	HPV18 L2	9	149
HPV18		11	286	HPV18 L2	8	377
HPV18		8	341	HPV18 L2	9	39
HPV18		11	341	HPV18 L2	9	406
HPV18		11	322	HPV18 L2	8	367
HPV18		11	404	HPV18 L2	9	114
HPV18		11	443	HPV18 L2	9	288
HPV18		11	241	HPV18 L2	9	392
HPV18		11	296	HPV18 L2	10	148
HPV18		8	429	HPV18 L2	10	38
HPV18		10 -	429	HPV18 L2	9	154
HPV18		10	62	HPV18 L2 HPV18 L2	9	366
HPV18		8	64	HPV18 L2 HPV18 L2	11	388 217
HPV18		10	432	HPV18 L2	10	339
HPV18		11	432 183	HPV18 L2	8	150
HPV18		10	310	HPV18 L2	11	417
HPV18 HPV18		10 11	310	HPV18 L2	8	234
HPV18		11	37	HPV18 L2	10	113
HPV18		10	44	HPV18 L2	9	387
HPV18		10	323	HPV18 L2	11	112
HPV18		11	152	HPV18 L2	9	427
HPV18		10	405	HPV18 L2	10	427
HPV18		8	249	HPV18 L2	8	436
HPV18		11	43	HPV18 L2	11	374
HPV18		9	248	HPV31 E1	10	186
HPV18		10	242	HPV31 E1	8	504
HPV18		10	287	HPV31 E1	9	81
HPV18		10	391	HPV31 E1	9	213
HPV18		11	338	HPV31 E1	8	96
HPV18	L2	10	386	HPV31 E1	8	421
HPV18	L2	8	325	HPV31 E1	8	336
HPV18	L2	11	390	HPV31 E1	10	336
HPV18	L2	8	362	HPV31 E1	9	364
HPV18	L2	10	362	HPV31 E1	8	352
HPV18		11	362	HPV31 E1	9	42
HPV18	L2	9	419	HPV31 E1	10	348
HPV18		9	120	HPV31 E1	9	311
HPV18		9	376	HPV31 E1		583
HPV18		8	185	HPV31 E1	11	583
HPV18		8	258	HPV31 E1	8	50
HPV18		10	360	HPV31 E1	9	473
HPV18		8	312	HPV31 E1	9	425
HPV18		9	312	HPV31 E1	8	436
HPV18		10	172	HPV31 E1	8	199
HPV18		9	233	HPV31 E1	9	566
HPV18		9	298	HPV31 E1	11	433
HPV18		9	268	HPV31 E1	11	499
HPV18		8	364	HPV31 E1 HPV31 E1	9	305 252
HPV18	L2	9	364	HAART ET	TO	252

Table VII HLA-A1 Supermotif Peptides

				•			
HPV31	E1	8	11	HPV31	E2	9	307
HPV31	E1	9	11 .	HPV31	E2	11	22
HPV31	E1	11	196	HPV31	E2	8	124
HPV31	E1	10	222	HPV31	E2	11	124
HPV31	E1	9	243	HPV31	E2	11	197
HPV31		8	328	HPV31	E2	8	80
HPV31		9	560	HPV31		11	185
HPV31		11	478	HPV31		8	200
HPV31		11	309	HPV31	E2	8	171
HPV31		11	471	HPV31	E2	11	168
HPV31		10	479		E2	10	35
HPV31		11	479	HPV31 HPV31		9	164 345
HPV31		10	291 291	HPV31	E2 E2	8	193
HPV31		11	590	HPV31	E2	8	312
HPV31 HPV31		8 11	463	HPV31	E2	10	78
HPV31		8	119	HPV31		11	77
HPV31		10	232		E2	8	303
HPV31		8	412	HPV31	E2	9	84
HPV31		8	234	HPV31	E2	8	127
HPV31		10	94	HPV31	E2	10	127
HPV31		9	584	HPV31	E2	9	361
HPV31	E1	10	584	HPV31	E2	11	9
HPV31	E1	9	337	HPV31	E2	10	106
HPV31	E1	10	563	HPV31	E2	10	317
HPV31	E1	10	500	HPV31	E2	10	191
HPV31	E1	9	187	HPV31	E2	8	151
HPV31		8	306		E2	9	151
HPV31		11	47	HPV31	E2	8	321
HPV31		9	253	HPV31		8	25
HPV31		10	547	HPV31		9	25
HPV31		10	117	HPV31 HPV31	E2	9	37 311
HPV31 HPV31		11	93 580	HPV31	E2	8	346
HPV31		10	207	HPV31		10	198
HPV31		8	323	HPV31		9	128
HPV31		10	124	HPV31		11	128
HPV31		11	562	HPV31		10	93
HPV31		8	293	HPV31	E2	8	362
HPV31		9	293	HPV31	E2	9	192
HPV31		10	293	HPV31	E2	11	92
HPV31	E1	11	303	HPV31	E2	10	344
HPV31	E1	11	40		E2	8	131
HPV31	E1	8	294	HPV31		9 .	159
HPV31		9	294	HPV31	E2	10	159
HPV31		11	211	HPV31	E5	11	40
HPV31		9	233		E5	8	53
HPV31		11	333		E5	11	53
HPV31		11	505	HPV31		8	61
HPV31		11	325		E5	10 9	15 72
HPV31		9	349	HPV31 HPV31	E5 E5	10	6
HPV31 HPV31		11	349 254		E5	9	11
HPV31		10	434		E5	9	16
HPV31		10	197	HPV31		8	43
HPV31		9	223		E5	9	42
HPV31		9	564		E5	8	32
HPV31		11	564		E5	11	5
HPV31		9	558	HPV31	E5	11	70
HPV31		11	558	HPV31	E5	8	56

Table VII HLA-A1 Supermotif Peptides

HPV31	B5	11	56	HPV31	L1	11	381
HPV31	E5	9	31	HPV31	L1	8	357
HPV31	E5	10	10	HPV31	L1	10	65
HPV31	E5	9	7	HPV31	L1	8	20
HPV31	E5	10	41	HPV31		8	42
HPV31	E5	10	54	HPV31		9	42
HPV31		8	8	HPV31		8	384
HPV31		10	51	HPV31		8	43
HPV31		8	73	HPV31		11	238
HPV31		8	12	HPV31		11	201
HPV31		11	9	HPV31 HPV31		9 11	300 351
HPV31		9	64 50	HPV31		9	227
HPV31		11 10	63	HPV31		11	222
HPV31 HPV31		11	66	HPV31		9	411
HPV31		8	63	HPV31		11	411
HPV31		8	25	HPV31		11	17
HPV31		10	14	HPV31		8	306
HPV31		9	39	HPV31	L1	8	255
HPV31		8	47	HPV31	L1	9	41
HPV31	E6	9	61	HPV31	L1	10	41
HPV31	E6	10	61	HPV31	L1	8	77
HPV31	E6	8	118	HPV31		10	77
HPV31		8	72	HPV31		10	75
HPV31		10	72	HPV31		8	228
HPV31		9	15	HPV31		8	414
HPV31		9	37	HPV31 HPV31		11	2 394
HPV31		11	37	HPV31			299
HPV31		10 8	36 16	HPV31			283
HPV31 HPV31		9	73	HPV31			283
HPV31		9	132	HPV31			286
HPV31		9	70	HPV31		9	383
HPV31		10	70	HPV31		11	302
HPV31		10	48	HPV31	L1	8	354
HPV31		8	4	HPV31	L1	11	354
HPV31	E7	10	78	HPV31		11	267
HPV31	E7	11	77	HPV31		9	66
HPV31	E7	9	49	HPV31			18
HPV31		9	348	HPV31		8	28
HPV31		8	398	HPV31			301 62
HPV31		8	285	HPV31 HPV31		10 9	235
HPV31 HPV31		9	285 224	HPV31			364
HPV31		11	459	HPV31		8	250
HPV31		8	129	HPV31			27
HPV31		9	203	HPV31		9	27
HPV31		9	353	HPV31	L2	11	286
HPV31		8	270	HPV31	L2	9	311
HPV31		8	449	HPV31	L2	10	311
HPV31	L1	8	456	HPV31	L2		311
HPV31	L1	9	323	HPV31	L2	11	376
HPV31	L1	11	117	HPV31			354
HPV31		9	413	HPV31			253
HPV31		11	298	HPV31		11	253
HPV31		11	282	HPV31		11	237
HPV31		11	393	HPV31 HPV31	L2	8 11	433 351
HPV31		10	118	HPV31		10	63
HPV31		10	382 61	HPV31		8	65
HPV31	ыl	11	9.1	nrv31		•	-

Table VII HLA-A1 Supermotif Peptides

HPV31	L2	11	213		HPV33	E1	8	349
HPV31		11	38		HPV33	E1	10	349
HPV31	L2	10	45		HPV33	E1	8	365
HPV31		11	45		HPV33	E1	9	42
HPV31		8	245		HPV33	E1	9	377
HPV31		9	244		HPV33	E1	9	62
HPV31		10	238		HPV33	El	9	324
HPV31		11	178		HPV33		9	516
HPV31		10	395		HPV33		10	361
HPV31		10	287		HPV33	E1	11	361
HPV31		11	447		HPV33	E1	8	449
HPV31		8	269		HPV33	E1	8	212
HPV31		10	3 90		HPV33	E1		446
HPV31		10	410		HPV33	E1	11	446
HPV31		11	122		HPV33	E1	10	265
HPV31		11	394		HPV33	E1	11	209
HPV31		9	425		HPV33	E1	8	11
HPV31		11	44		HPV33	E1	11	512
HPV31		10	243		HPV33 HPV33	E1 E1	8	564 341
HPV31		9	378		HPV33	E1	9	573
HPV31		9	229 429		HPV33	E1	11	192
HPV31		11	9		HPV33	E1	9	266
HPV31 HPV31		9	431		HPV33	E1	8	267
HPV31		10	431		HPV33	E1	9	200
HPV31		8	181		HPV33	E1	10	492
HPV31		9	180		HPV33	E1	11	492
HPV31		10	179		HPV33	E1	11	322
HPV31		9	396		HPV33	E1	10	210
HPV31		8	151		HPV33	E1	9	520
HPV31		8	346		HPV33	E1	10	124
HPV31		11	346		HPV33	E1	10	304
HPV31		8	379		HPV33	E1	11	304
HPV31		10	149		HPV33	E1	10	220
HPV31		9	40		HPV33	E1	8	603
HPV31	L2	8	312		HPV33	E1	11	476
HPV31	L2	9	312		HPV33	E1	8	425
HPV31	L2	10	312		HPV33	E1	10	245
HPV31		10	347		HPV33	E1	8	247
HPV31	L2	11	266		HPV33	E1	9	438
HPV31	L2	9	288		HPV33	E1	9	350
HPV31	L2	9	345		HPV33	E1	9	362
HPV31	L2	11	148		HPV33	E1	10	362
HPV31		10	39		HPV33	E1	11	362
HPV31	L2	8	426		HPV33	E1	10	576
HPV31		10	344		HPV33	E1	8	336
HPV31		11	343		HPV33	El	10	513
HPV31		9	391		HPV33	E1	11	443
HPV31		9	254		HPV33	E1	11	346
HPV31		10	254		HPV33	E1	10	199
HPV31		8	392		HPV33	E1	8	195
HPV31		10	430		HPV33	E1	10	195
HPV31		11	430		HPV33	E1	10	560
HPV31		9	150		HPV33	E1	10	519 434
HPV33		10	596		HPV33 HPV33	E1 E1	8 10	593
HPV33		11	596					
HPV33		9	81		HPV33	E1 E1	10	437 308
HPV33		9	226		HPV33	E1	11	575
HPV33		8	494		HPV33	E1	9	335
HPV33	El	9	494		ubany	PT	,	232

Table VII HLA-A1 Supermotif Peptides

HPV33	E1	8	306	н	V33	E2	8	131
HPV33		9	306		V33		11	56
HPV33		10	306		V33		10	3
HPV33		10	111		V33		9	42
HPV33		10	193		V33		10	42
HPV33		11	224		V33		8	5
HPV33		11	110		V33		8	44
		9	577		V33		10	44
HPV33			577		V33		9	23
HPV33		11	491		V33		9	48
HPV33			246		V33		11	48
HPV33		9			V33		8	22
HPV33		11	338		V33		10	22
HPV33		8	517 571		V33		9	32
HPV33		9			V33		11	32
HPV33		11	571					
HPV33		10	78		V33		8	24
HPV33		11	41		V3 3		8	35
HPV33		10	10		V3 3		8	33
HPV33		9	288		V33		10	33
HPV33		10	145			E5	9	1
HPV33		9	25			E5	9	21
HPV33		10	235		V3 3		11	21
HPV33		10	298		V3 3		8	46
HPV33		10	282		V33	E5	11	46
HPV33		8	80		V33		9	34
HPV33		11	100		V33		10	31
HPV33		10	325		V33		11	40
HPV33		11	34		V33		9	58
HPV33		9	84		V33		11	66
HPV33		11	23		V33		8	69
HPV33		8	151		V33		11	69
HPV33		9	151		V33		9	61
HPV33		10	35		V33		8	118
HPV33		9	62		V33		9	73
HPV33		10	42		V33		8	72
HPV33		11	82		V33		10	72
HPV33		8	147		V33		10	70
HPV33		11	315		V33		11	50
HPV33	E2	8	284		V33		8	36
HPV33	E2	8	127		V33		10	36
HPV33		11	60		V33		9	39
HPV33		9	342		V33		10	51
HPV33		9	292		V33		9	52
HPV33		8	37		V33		10	14
HPV33		10	61		V33		11	6
HPV33	E2	8	302		V3 3		10	7
HPV33		9	301		V33		10	392
HPV33		10	93		V33		8	284
HPV33		11	128		V33		9	284
HPV33		9	146		V33		9	411
HPV33		8	343		V33		10	345
HPV33		9	326		V33		9	223
HPV33	E2	11	148		V33		8	396
HPV33		9	102		V33		11	457
HPV33		11	92		V33		9	351
HPV33		9	159		V33		8	129
HPV33		10	159		V33		9	202
HPV33		8	300		V33		9	303
HPV33		10	300		V33		10	303
HPV33	E2	8	44	HP	V33	L1	8	447

Table VII HLA-A1 Supermotif Peptides

HPV33	Ll	8	249	HPV33	L1	10	27
HPV33	L1	8	454	HPV33	L2	11	291
HPV33	L1	9	322	HPV33	L2	10	272
HPV33	L1	11	117	HPV33	L2	10	431
HPV33	L1	11	297	HPV33	L2	11	258
HPV33	L1	9	226	HPV33	L2	10	447
HPV33	L1	11	281	HPV33	L2	11	242
HPV33	L1	9	365	HPV33	L2	11	183
HPV33		11	365	HPV33	L2	8	440
HPV33		10	118	HPV33	L2	8	421
HPV33	L1	10	65	HPV33	L2	10	421
HPV33	L1	11	379	HPV33	L2	8	64
HPV33		8	20 .	HPV33	L2	10	62
HPV33		8	42	HPV33	L2	11	218
HPV33		9	42	HPV33	L2	11	37
HPV33		11	61	HPV33	L2	8	374
HPV33		8	382	HPV33	L2	11	374
HPV33		10	62	HPV33	L2	8	336
HPV33		11	237	HPV33	L2	10	44
HPV33		11	200	HPV33	L2	11	44
HPV33		9	299	HPV33	L2	9	448
HPV33		11	221	HPV33	L2	11	448
HPV33		8	439	HPV33		9	273
HPV33		9	409	HPV33		9	155
HPV33		11	409	HPV33		10	292
HPV33		11	17	HPV33		8	250
HPV33		8	305	HPV33		11	250
HPV33		8	254	HPV33		10	104
HPV33		8	347	HPV33		8	433
HPV33		9	41	HPV33	L2	10	248
HPV33		10	41	HPV33	L2	9	249
HPV33		8	77	HPV33		10	243
HPV33		10	77	HPV33		11	405
HPV33		10	75	HPV33		10	372
HPV33		8	285	HPV33		10	391
HPV33		8	412	HPV33		8	423
HPV33		10	298	HPV33		11	333
HPV33		11	39	HPV33		9	413
HPV33		8	227	HPV33		10	347
HPV33		8	352	HPV33	L2	9	376
HPV33		11	352	HPV33	L2	9	121
HPV33		11	2	HPV33		11	411
HPV33		11	266	HPV33	L2	8	186
HPV33		9	381	HPV33	L2	8 '	221
HPV33		11	349	HPV33	L2	8	317
HPV33		10	238	HPV33	L2	9	317
HPV33		11	238	HPV33	L2	11	43
HPV33		11	301	HPV33	L2	11	191
HPV33		10	282	. HbA33	L2	11	153
HPV33		11	282	HPV33	L2	9	234
HPV33		9	66	HPV3 3	L2	10	357
HPV33		10	18	HPV33	L2	8	393
HPV33		8	28	HPV33		8	122
HPV33		9	28	HPV33		11	103
HPV33		10	380	HPV33		8	106
HPV33		8	300	HPV33		10	418
HPV33		8	362	HPV33		11	418
HPV33		9	234	HPV33		10	184
HPV33		8	27	HPV33		10	354
HPV33		9	27	HPV33		8	156
E v 3 3		-					

Table VII HLA-A1 Supermotif Peptides

				TIDIT III Dapellion	replaces			
HPV33	1.2	10	38		HPV45	E1	11	476
HPV33		9	39		HPV45		10	245
HPV33		10	154		HPV45		8	247
HPV33		9	432		HPV45		8	267
HPV33		9	244		HPV45		9	350
HPV33		9	293		HPV45		10	210
HPV33	L2	11	417		HPV45	E1	9	362
HPV33		11	353		HPV45		11	362
HPV33	L2	9	392		HPV45		9	200
HPV33	L2	9	105		HPV45	E1	10	513
HPV33	L2	8	356		HPV45	E1	10	560
HPV33	L2	11	356		HPV45	E1	8	414
HPV45	E1 ·	10	199		HPV45	E1	11	473
HPV45	E1	11	512		HPV45	E1	8	434
HPV45	E1	11	40		HPV45	E1	10	593
HPV45	E1	8	517		HPV45	E1	10	412
HPV45	E1	9	517		HPV45	E1	10	80
HPV45	E1	11	202		HPV45	E1	10	128
HPV45	E1	10	423		HPV45	E1	8	306
HPV45	E1	9	226		HPV45		9	306
HPV45		8	349		HPV45		10	306
HPV45	E1	10	349		HPV45	E1	11	575
HPV45	E1	10	361		HPV45	E1	8	307
HPV45	E1	10	623		HPV45		9	307
HPV45	E1	9	42		HPV45		8	308
HPV45	E1	10	596		HPV45		9	246
HPV45		11	596		HPV45		11	224
HPV45		8	365		HPV45		9	577
HPV45	E1	9	573		HPV45		11	577
HPV45		9	324		HPV45		9	81
HPV45		11	446		HPV45		9	266
HPV45		10	385		HPV45		8	325
HPV45		9	486		HPV45		10	576
HPV45		8	449		HPV45		9	571
HPV45		9	438		HPV45		11	571
HPV45		8	212		HPV45		10	84
HPV45		9	579		HPV45		10	16
HPV45		8	130		HPV45		9	305
HPV45		8	494		HPV45		11 8	134
HPV45		9	494		HPV45		8 9	158 158
HPV45 HPV45		11 8	209 11		HPV45		8	31
HPV45		9	11		HPV45		9	31
HPV45		11	459		HPV45		11:	28
HPV45		10	265		HPV45		8	171
HPV45		10	235		HPV45		8	319
HPV45		9	256		HPV45		11	106
HPV45		10	519		HPV45		8	154
HPV45		11	338		HPV45		9	154
HPV45		11	491		HPV45		10	41
HPV45		11	322		HPV45		9	341
HPV45		10	447		HPV45		8	301
HPV45		10	492		HPV45		9	187
HPV45		11	492		HPV45		9	357
HPV45		8	425		HPV45		8	109
HPV45		10	304		HPV45		9	332
HPV45		11	304		HPV45		11	40
HPV45		10	220		HPV45		9	90
HPV45		8	387		HPV45		8	43
HPV45		8	476		HPV45	E2	9	309

Table VII HLA-A1 Supermotif Peptides

				men-rer supermon	repudes			
HPV45	E2	11	142		HPV45	L1	8	111
HPV45		8	358		HPV45		8	143
HPV45		10	99		HPV45	Li	11	326
HPV45		8	159		HPV45	Li	11	422
HPV45		8	138		HPV45		9	396
HPV45		11	98		HPV45	L1	11	396
HPV45		9	166		HPV45	L1	10	12
HPV45		10	166		HPV45	Li	11	11
HPV45		8	145		HPV45	Li	10	5
HPV45		8	317		HPV45		10	411
HPV45		10	317		HPV45	L1	9	328
HPV45		11	175		HPV45		10	91
HPV45		8	137		HPV45	L1	8	68
HPV45		9	137		HPV45		9	68
HPV45		9	37		HPV45		8	413
HPV45		11	37		HPV45		8	69
HPV45		8	27		HPV45	L1	11	264
HPV45		10	77		HPV45	L1	11	227
HPV45		11	43		HPV45		11	4
HPV45		10	53		HPV45	L1	11	310
HPV45		8	120		HPV45	L1 .	8	425
HPV45		9	54		HPV45	L1	10	425
HPV45		8	92		HPV45	L1	8	383
HPV45		8	74		HPV45	L1	11	383
HPV45		9	41		HPV45	L1	8	17
HPV45		11	24		HPV45		9	22
HPV45		11	89		HPV45	L1	11	248
HPV45		8	38		HPV45	L1	11	139
HPV45		10	38		HPV45	L1	وَ	440
HPV45	E6	9	72		HPV45	L1	11	440
HPV45	E6	10	72		HPV45	L1	11	380
HPV45	E7	9	83		HPV45	L1	8	281
HPV45	E7	10	20		HPV45	L1	8	334
HPV45	E7	11	91	•	HPV45	L1	9	253
HPV45	E7	10	92		HPV45		9	67
HPV45	E7	9	89		HPV45	L1	10	67
HPV45	E7	9	93		HPV45		10	101
HPV45	E7	8	94		HPV45		8	46
HPV45	L1	8	103		HPV45		8	254
HPV45	L1	11	28		HPV45		8	427
HPV45		11	375		HPV45	L1	10	327
HPV45		8	88		HPV45	L1	8	443
HPV45		10	88			L1	10	423
HPV45		8	276			Ll	9	426
HPV45		9	188		HPV45	L1	11	65
HPV45		9	250		HPV45	L1	10	376
HPV45		11	488		HPV45		11	43
HPV45		9	332		HPV45	L1	11	412
HPV45		10	332		HPV45	L1	9	330
	L1	9	229		HPV45	L1	11	442
HPV45		11	461		HPV45	L1 L1	9 10	442
HPV45		8	296		HPV45	L1	8	329
HPV45		10	169		HPV45	L1	8	441
HPV45		8	313		HPV45	L1	10	441
	Ll	8	14		HPV45	L1	8	478
HPV45		11	14 485		HPV45	L1	11	293
HPV45		8			HPV45	L1	9	92
HPV45	L1	9	351 141		HPV45	ь1	8	54
HPV45 HPV45		10	141		HPV45	L1	9	477
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Table VII HLA-A1 Supermotif Peptides

				ALEX - A COOPERING	i i epiides			
HPV45	L1	9	261		HPV45	L2	9	154
HPV45	Ll	8	393		HPV45	L2	11	358
HPV45	Ll	8	53		HPV45	L2	9	149
HPV45	L1	9	53		HPV45	L2	8	363
HPV45	L2	11	286		HPV45	L2	11	363
HPV45	L2	9	114		HPV45	L2	9	39
HPV45	L2	8	340	•	HPV45	L2	10	376
HPV45	L2	11	340		HPV45	L2	9	393
HPV45	L2	11	405		HPV45	L2	8	155
HPV45	L2	9	345		HPV45	L2	9	268
HPV45	L2	8	343		HPV45	L2	11	418
HPV45	L2	11	343		HPV45	L2	10	38
HPV45	L2	10	148		HPV45	L2	10	359
HPV45		11	241		HPV45	L2	11	426
HPV45		11	296		HPV45	L2	8	389
HPV45		8	430		HPV45	L2	11	217
HPV45		10	430		HPV45	L2	8	150
HPV45		8	64		HPV45	L2	8	249
HPV45		10	62		HPV45	L2	9	388
HPV45		10	183		HPV45	L2	11	112
HPV45	L2	9	433		HPV45	L2	9	428
HPV45	L2	10	433		HPV45	L2	10	428
HPV45		11	433		HPV45	L2	8	437
HPV45		11	37		HPV45	L2	9	437
	L2	10	406		HPV56	E2	10	21
HPV45		9	407		HPV56	E2	9	71
HPV45	L2	10	44		HPV56	E2	11	71
HPV45		10	338		HPV56	E2	10	92
HPV45		11	152		HPV56	E2	11	92
HPV45		11	43		HPV56	E2	9	140
HPV45	L2	8	366		HPV56	E2	8	263
	L2	11	337		HPV56	E2	11	43
HPV45		10	287		HPV56	E2	8	23
HPV45	L2	10	242		HPV56	E2	10	128
HPV45	L2	11	375		HPV56	E2	11	294
	L2	10	392		HPV56	E2	8	261
HPV45		9	248		HPV56	E2	10	261
HPV45	L2	10	387		HPV56	E2 E2	9	66 94
HPV45		8	258		HPV56 HPV56	E2	9	94
HPV45	L2	11	391		HPV56	E2 E2	8	130
	L2	8	378		HPV56	E2	8	297
HPV45		8	361 361		HPV56	E2	10	299
HPV45 HPV45	L2	10 9	120		HPV56	E2	11	258
HPV45	L2	9	420		HPV56	E2	8	90
	L2	8	185		HPV56	E2	10	295
HPV45		10	267		HPV56	E2	11	25
	L2	11	118		HPV56	E2	8	46
HPV45	L2	8	312		HPV56	E2	11	149
HPV45	L2	9	312		HPV56	E2	8	152
HPV45		10	172		HPV56	E2	8	301
HPV45	L2	9	233		HPV56	E2	9	246
HPV45	L2	10	451		HPV56	E2	10	26
HPV45	L2	9	298		HPV56	E2	8	141
HPV45	L2	8	220		HPV56	E2	8	28
	L2	10	247		HPV56	E2	10	259
HPV45	L2	11	246		HPV56	E2	10	36
HPV45	L2	9	288		HPV56	E2	8	271
HPV45	L2	10	153		HPV56	E2	9	27
	L2	9	362		HPV56	E2	10	150
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Table VII HLA-A1 Supermotif Peptides

			•				
HPV56	E2	9	45	HPV56	L1	8	334
HPV56		11	35	HPV56		8	258
HPV56		9	270	HPV56	L1	11	258
HPV56	E2	10	79	HPV56	L1	11	413
HPV56	E2	8	278	HPV56	L1	10	93
HPV56		11	111	HPV56	L1	11	300
HPV56	E2	8	74	HPV56	L1	10	98
HPV56		9	74	HPV56	Ll	8	55
HPV56	E2	9	102	HPV56	L1	8	77
HPV56	E2	10	102	HPV56	L1	9	77
HPV56	E6	9	64	HPV56	L1	8	416
HPV56	E6	10	64	HPV56	L1	11	234
HPV56	E6	11	69	HPV56	L1	9	333
HPV56	E6	8	50	HPV56	L1	8	2
HPV56	E6	8	28	HPV56		9	1
HPV56	E6	10	52	HPV56	L1	11	271
HPV56	E6	8	39	HPV56	L1	8	95
HPV56	E6	10	39 .	HPV56	L1	10	95
HPV56	E6	8	54	HPV56		10	426
HPV56	E6	10	54	HPV56		9	31
HPV56	E6	8	75	HPV56	L1	8	473
HPV56	E6	10	75	HPV56	L1	11	255
HPV56	E6	10	26	HPV56	L1	9	13
HPV56	E6	10	70	HPV56		11	467
HPV56	E6	9	40	HPV56		10	442
HPV56	E6	9	55	HPV56	L1	11	52
HPV56	E6	11	25	HPV56		8	288
HPV56	E6	10	98	HPV56		8	339
HPV56	E6	10	119	HPV56		9	260
HPV56	E6	9	135	HPV56		9	76
HPV56	E6	9	73	HPV56		10	76
HPV56	E6	10	73	HPV56		8	110
HPV56		9	62	HPV56		10	108
HPV56	E7	11	60	HPV56		8	446
HPV56		8	381	HPV56		10	332
HPV56		8	444	HPV56		11	74
HPV56		10	444	HPV56		10	379
HPV56		11	37	HPV56		8	261
HPV56		8	26	HPV56		9	415
HPV56		9	195	HPV56		11	335
HPV56		9	257	HPV56		9	445
HPV56		11	491	HPV56		9	99
HPV56		10	486	HPV56		10	53
HPV56		10	60	HPV56		10	7 268
HPV56		11	60	HPV56		9	396
HPV56		9	236	HPV56		8	283
HPV56		8	23	HPV56 HPV56		8	62
HPV56		11	23	HPV56		9	62
HPV56		8	481	HPV56		8	438
HPV56		8	303	HPV56		11	246
HPV56		10	21	HPV56		11	406
HPV56		9	488	HPV56		11	30
HPV56		-	356	HPV56		9	429
HPV56		8	118	HPV56		9	114
HPV56		8	150	HPV56		10	287
HPV56		11	331	HPV56		11	118
HPV56		9	399	HPV56		8	64
HPV56		11	399	HPV56		10	434
HPV56		11	378	HPV56		11	434
HPV56	ьī	10	414	nrvso			. 34

Table VII HLA-A1 Supermotif Peptides

SF 1168080 vl

HPV56 L2 8 258 HPV56 L2 10 62 HPV56 L2 10 310 HPV56 L2 11 310 HPV56 L2 8 269 11 11 8 372 HPV56 L2 11 190 HPV56 L2 HPV56 L2 44 HPV56 L2 10 44 HPV56 L2 11 44 HPV56 L2 9 210 HPV56 L2 11 182 HPV56 L2 9 279 HPV56 L2 10 407 HPV56 L2 9 43 HPV56 L2 11 43 HPV56 L2 10 38 HPV56 L2 8 337 338 HPV56 L2 11 248 HPV56 L2 9 278 HPV56 L2 10 HPV56 L2 10 342 HPV56 L2 10 388 395 HPV56 L2 8 HPV56 L2 9 374 HPV56 L2 10 209 HPV56 L2 10 392 HPV56 L2 11 392 HPV56 L2 9 336 HPV56 L2 10 267 HPV56 L2 9 410 HPV56 L2 8 185 HPV56 L2 8 312 HPV56 L2 9 312 HPV56 L2 10 312 421 HPV56 L2 11 HPV56 L2 9 233 220 HPV56 L2 8 HPV56 L2 9 435 HPV56 L2 10 435 HPV56 L2 11 435 HPV56 L2 10 153 HPV56 L2 8 211 HPV56 L2 9 154 HPV56 L2 10 183 HPV56 L2 11 414 HPV56 L2 10 247 HPV56 L2 9 288 HPV56 L2 11 112 HPV56 L2 9 408 HPV56 L2 11 408 HPV56 I.2 8 249 HPV56 L2 11 152 HPV56 L2 9 389 HPV56 L2 10 31 HPV56 L2 10 431

116

Table VIIA HPV6A HLA-A1 Supermotif Peptides

2	3	4		L1	8	450
L2	11	286		E1	9	587
E4	8	14		E2	8	171
E1	11	520		E5	9	28
E1.	10	207		L1	9	318
L1	8	81		E1	10	243
L2	8	421		E2	9	156
E6	8	37		E1	11	217
E6	10	37		E1	10	273
L2	9	288		E1	8	11
E1.	11	330		L2	10	431
L1	9	342		L2	11	431
E1	8	525		L2	10	62
E6	11	10		E1	10	431
E1	10	77		L1	11	293
E1	10	601		E2	10	179
E6	11	67		L2	11	215
E2	10	35		L2	8	64
E6	11	131		E1	11	436
E4	9	64		L1	9	407
E1	10	369		Li	9	222
	9	219		E1	8	316
L1		96		Ll	9	111
E6	10			Li	11	113
E1	8	570		L2	8	312
E1	10	570		L2	9	312
E2	9	313		L2	10	312
E1	11	81				
E2	9	25		E6	8	119
E1	10	203		E1	9	264
E1	9	42		E4	8	59
L2	11	266		E2	10	78
E7	9	44		E2	10	310
L2	10	344		E4	10	10
L1	9	198		E2	9	338
E2	10	136		E2	10	149
L2	10	120		E2	11	149
L1	11	453		E6	9	25
E1	10	604		L1	11	387
E1	11	604		E1	9	581
E1	9	131		L1	9	361
E1	10	417		L1	11	361
E2	11	100		E1	8	502
E1	8	373		E1	9	502
E2	8	80		E5	8	21
E2	8	293		E5	10	31
E2	10	293		E6	9	97
L1	8	443		E5	9	32
E2	10	205		L2	10	44
E1	8	220		L2	11	44
E6	8	126		E5	8	17
E1	8	454		E1	10	500
E1	11	454		E1	11	500
L2	8	428		E1	9	571
E5	9	68		L1	10	376
E1	10	393		L2	8	247
L2	10	398		E1	11	476
E1	9	446		L2	9	121
L1	8	245		E5	10	34
	8	457		E1	8	433
E1		239		E6	8	73
L2	11	- 37			-	

Table VIIA HPV6A HLA-A1 Supermotif Peptides

E6	10	73	E2	10	281
E1	10	312	L2	10	38
E1	11	312	E2	8	127
E1	9	254	L1	11	217
E1	8	357	L2	10	189
E1	10	357	L1	11	109
E1	10	228	E1	10	258
E1	11	484	L2	10	389
E2	9	84	L2	10 '	337
L1	10	56	L1	9	391
E6	11	116	L2	9	408
E6	8	52	E2	9	354
E6	10	52	L1	11	426
L1	10	61	L1	8	90
Li	8	19	L2	9	426
Li	10	71	L2	10	426
E1	8	255	E5	10	19
E5	و	16	L1	11	16
E2	8	314	L2	11	418
L2	9	246	L2	9	363
E5	8	33	L2	11	43
E5	11	33	L1	8	301
L1	8	41	E6	10	50
L1	9	41 -	E4	9	4
E1	10	521	L1	8	250
E1	9	208	L2	8	400
E2	11	82	E1	8	314
E5	9	59	E1	9	314
E5	10	59	E1	10	314
E5	8	51	E2	8	103
E5	8	69	E1	10	128
E5	8	60	L2	9	231
E5	9	60	L2	10	245
E5	10	72	L1	9	40
L1	8	378	L1	10	40
E1	10	218	E2	10	303
E1	9	259	L1	9	279
E1	9	605	Li	8	140
	10	605	E1	11	583
E1	9	390	L2	8	153
L2 L2	10	240	L2	10	267
E1		132	E5	11	30
	8 9	358	E2	8	207
E1	10	49	L1	11	375
E5 E6	9	38	E1	8	60
E6	11	38	L1	10	294
		58 61	El	8	260
E5	8 9		E6	11	23
L2		338	E2	9	150
E5	9	73	E2	10	150
E5	8	47	E2	9	282
L1	9	295	L1	11	297
E2	8	151	L2	8	391
E2	9	151		11	
L1	11	196	L1	9	38 347
E1	9	274	L1	11	23
E1	10	568	E2		
E1 -	11	451	E2	9	180
E4	10	57	E5	11	14
E1	9	59	L2	10 9	374
E1	8	395	L2	9	241

Table VIIA HPV6A HLA-A1 Supermotif Peptides

E1	10	331
L1	8	348
L1	11	348
L1	8	392
E5	10	4.5
L2	11	145
L1	8	343
L2	9	39
E6	9	12
E2	8	355
L1	8	408
E5	10	15
E5	9	50
E5	11	71
E2	10	93
E6	8	26
L1	9	377
E2	11	128
	10	388
Ll	9	147
L2		
L2	9	152
Ll	10	346
L2	11	373
L1	8	280
E5	11	44
E6	8	39
E6	10	39
E1	11	232
L1	8	223
E1	9	585
E1	11	585
		11
E6	10	
L2	10	151
L1		345
L2	11	150
E1	9	332
E1	9	78
L1	9	57
L1	11	57
E1	11	346
E1	8	333
E5	9	20
E1	11	499
E6	9	53
E1	8	275
	8	73
L1		
E5	11	48
E5	9	46
L1	10	114
L1	9	62
E5	8	29
E2	9	206
L1	10	17
Ll	8	296
L2	10	419
L2	8	364
L1	8	27
		146
L2	10	584
E1	10	
L1	9	72

Table VIIB HPV6B HLA-A1 Supermotif Peptides

		· ·	•		
2	3	4	L2	11	239
L2	11	286	E1	8	457
E4	8	24	Ll	8	450
E1	11	520	El	9	587
E1	10	207	E2	8	171
L1	8	81	E5A	9	28
L2	8	421	L1	9	318
E6	8	37	E1	10	243
E6	10	37	E2	9	156
L2	9	288	E1	11	217
E1	11	330	E5B	8	15
L1	9	342	E5B	10	25
E1	8	525	E1	10	273
Eб	11	10	E1	8	11
E1	10	77	L2	10	431
E1	10	601	L2	11	431
E1	9	234	L2	10	62
E6	11	67	E1	10	431
E2	10	35	L1	11	293
E6	11	131	E2	10	179
E4	9	74	L2	11	215
E1	10	369	L2	8	64
L1	9	219	E1	11	436
E6	10	96	L1	9	407
E1	8	570	L1	9	222 316
E1	10	570 25	E1 L1	9	111
E2	9		L1	11	113
E2	9	313 81	L2	8	312
E1 E1	11	203	L2 L2	9	312
E2	9	338	L2	10	312
E1	9	42	E6	8	119
L2	11	266	E1	9	264
E7	9	44	E2	10	78
L2	10	344	E2	10	310
L1	9	198	E6	10	50
E2	10	136	E4	10	20
L2	10	120	E2	10	149
Li	11	453	E2	11	149
E1	10	604	E6	9	25
E1	11	604	L1	11	387
E1	9	131	E1	9	581
E1	10	417	L1	9	361
E2	11	100	Ll	11	361
E1	8	373	E1	8	502
E2	8	80	E1	9	502
E2	8	293	E5A	8	21
E2	10	293	E5A	10	31
L1	8	443	E6	9	97
E2	10	205	E5A	9	32
E1	8	220	L2	10	44
E6	8	126	L2	11	44
ESA	9	16	E5A	8	17
E1	8	454	E1	10	500
E1	11	454	El	11	500
L2	8	428	E1	9	571
E5A	9	68	L1	10	376
E1	10	393	L2	8	247
L2	10	397	El	11	476
E1	9	446	L2	9	121
L1	8	245	E5A	10	34

Table VIIB HPV6B HLA-A1 Supermotif Peptides

		HLA-A1 Supermoni i	epuaes		
E1	8	433	E4	10	67
E6	8	73	E2	10	281
E6	10	73	L2	10	38
		312	E2	8	127
	10		E1	8	607
E1	11	312			
E1	9	254	L1	11	217
E1	8	357	L2	10	189
E1	10	357	E4	8	69
E1	10	228	L1	11	109
E1 .	11	484	L2	9	389
L1	10	56	E1	10	258
E6	11	116	L2	10	337
E6	8	52	Ll	9	391
E6	10	52	L2	9	407
L1	10	51	E2	9	354
L1	8	19	L1	11	426
L1	10	71	L1	8	90
E1	8	255	L2	9	426
E2	8	314	L2	10	426
L2	9	246	E5A	10	19
E5A	8	33	L1	11	16
E5A	11	33	L2	9	363
L1	8	41	E5A	10	7
L1.	9	41	L2	11	43
E1	10	521	L1	8	301
E1	9	208	E4	9	14
E5A	9	59	L1	8	250
E5A	10	59	L2	8	399
E5A	8	51	E1	8	314
E2	11	82	E1	9	314
E5A	8	69	E1	10	314
E5A	8	60	E2	8	103
E5A	9	60	E1	10	128
E5A	10	72	L2	9	231
L1	8	378	L2	10	245
E1	10	218	Li	9	40
L2	8	390	Li	10	40
	9	259	E2	10	303
E1	9	605	E2	9	84
E1			L1	9	279
E1	10	605	L1	8	140
L2	10	240			583
E5B	11	3	E1 L2	11 8	153
E1	8	132			
E1	9	358	L2 E5A	10 11	267 30
E5A	10	49	E2A		207
E6	9	38			
E6	11	38	L1	11	375
E5A	8	61	E1	8	60
L2	9	338	L1	10	294
E5A	9	73	E1	8	260
E5A	8	47		11	23
L1	9	295	E2	9	150
E5B	9	26	E2	10	150
E2	8	151	E2		282
E2	9	151	L1	11	297
Ll	11	196	L1	11	38
E1	9	274			347
E1	10	568	E2	11	23
E1	11	451	E5A	11	14
E1	9	59		9	180
E1	8	395	L2	10	374

Table VIIB HPV6B HLA-A1 Supermotif Peptides

			HLA-
L2	9	241	
E1	10	331	
Li	8	348	
L1	11	348	
Li	8	392	
E5A	10	4.5	
L2	11	145	
L1	8 .	343	
L2	9	39	
E6	9	12	
E2	8	355	
L1	8	408	
ESA	9	50	
E5A	11	71	
E2	10	93	
E6	8	26	
L1	9	377	
E2	11	128	
ESB	10	59	
L1	10	388	
L2	9	147	
L2		152	
L1 L2	10 11	346 373	
L1	8	280	
ESA	11	44	
E6	8	39	
E6	10	39	
L1	8	223	
E1	11	232	
E5B	8	63	
E1	9	585	
E1	11	585	
E6	10	11	
L2	10	151	
L1	11	345	
L2 E1	11 9	150 . 332	
L2	11	387	
E1	9	78	
Ll	9	57	
Ll	11	57	
E1	11	346	
E1	8	333	
E5A	9	20	
El	11	499	
E6	9	53	
El	8	275	
Ll	8	73	
E5A	11	48	
E5A	9	46	
L1 L1	10 9	114 62	
E5A	8	29	
E2	9	206	
L1	10	17	
L1	8	296	
E5B	11	58	
L2	8	364	
L2	11	418	
L1	8	27	

E2 8 131

Table VIIC. HPV11 HLA-A1 Supermotif Peptides

				-		
2	. 3	_4		E5	8	16
L2	11	285		L2	11	295
E1	11	520		Ll	8	451
L1	8	81		E1	9	587
L2	8	417		E1	8	220
E6	8	37		L2	10	393
E6	10	37		L1 ·		319
E1	11	330		E1	10	243
L1	9	343		E2	9	156
E1	8	525		E1	11	217
E6	11	10		E1	10	273
E1	10	77		L2	10	427
L1	8	349		L2	11	427
L1	11	349		E1	8	11
E5	10	26		L2	8	63
E1	10	601		L1	11	294
E6	11	67		E2	10	179
E5	8	73		E1	10	431
E5	9	73		L1	9	408
E4	9	73		L1	9	223
E1	10	369		E1	8	316
L1	9	220		L1	11	113
E2	9	25		L2	8	311
E6	10	96		L2	9	311
E1	10	203		E6	8	119
E1	8	570		E1	9	264
E1	10	570		E2	10	136
E1	11	81		E2	10	78
E1	9	42		E2	- 10	309
E2	8	292		E5	8	7
E2	10	292		E5	10	7
L2	10	343		E4	10	20
L1	9	199		E1	8	349
E5	9	12		E1	9	581
L1	8	125		E6	9	25
E2 .	9	312		L1	11	388
Li	11	454		L2	11	36
E6	9	69		L2	10	188
E1	10	604		L1	9	362
E1	11	604		L1	11	362
E1	9	131		E5	10	34
E1	10	417		E5	9	35
E2	11	100		E6	9	97
E1	8	373		L2	10	43
L2	11	265		L2	11	43
E2	8	80		E5	8	17
E1	10	128		E1	9	571
L1	8	444		E1	10	500
E2	10	205		E1	11	500
L2	10	119		Ll	10	377
E6	8	126		L2	10	286
E1	11	454		E5	10	31
L2	8	424		E6	8	73
E1	9	494		E6	10	73
E5	9	68		E1	10	312
E5	10	68		E1	11	312
	10	393		E1	9	254
E1	9	446		E6	11	116
E1		445		E1	8	357
E1	8			E1	10	357
L2	11	238			10	33,

Table VIIC. HPV11 HLA-A1 Supermotif Peptides

			HLA-A i Supermotti	Peptides		
E1	10	420		E5	8	54
L1	10	347		E2	8	127
E1	11	484		L1		218
E1	10	228		L2	8	385
E5	10	50		L2	9	385
E2	9	84		E1	10	258
Ll	10	56		E4	8	68
E1	8	433		L2	10	336
L1	10	61		L1	9	392
L1	8	19		L2	9	403
L1	10	71		E2	9	353
E6	8	52		L1	11	427
E6	10	52		L2	11	206
E1	8	255		L1	8	90
E5	8	33		L2	9	422
E5	11	33		L2	10	422
E5	9	16		L2	10	358
E5	8	36		L1	11	16
L1	8	41		L2	11	42
L1	9	41		L1	8	302
E1	10	521		E4	9	14
E4	9	18		L1	8	251
E5	9	59		L2	8	395
E5	10	59		E1	8	314
E5	8	51		E1	9	314
E5	8	69		E1	10	314
E5	9	69		L2	9	230
E5	8	60		L2	9	297
E5	9	60	*	L1	9	40
L1	8	379		L1	10	40
E1	9	605		E2	10	302
E1	10	605		L2	10	244
E1	10	218		L1 E1	9 8	280
L2	8	386		Ll	8	205 141
E1	9	259 239		E1	11	583
L2	10 11	4		E2	8	207
E5 E1	8	132		E2	11	23
E1	9	358		E6	9	12
E5	8	70		L2	10	266
E5	11	70		E1	8	422
E5	10	49		L1	11	376
E2	В	103		E5	11	30
E6	9	38		E6	11	23
E6	11	38		L1	10	295
E5	8	61		E1	8	260
L2	9	337		L1	11	298
E5	8	47		E2	9	337
L1	9	296		Ll	11	38
L2	9	245		L2	9	208
E5	9	62		L1	8	281
E5	11	62		E2	9	150
E1	9	421		E2	10	150
L1	1.1	197		E2	11	260
E1	9	274		E1	11	206
E4	9	1		E2	9	180
E1	10	568		L2	8	209
E6	10	50		L2	10	370
E1	8	395		L2	9	240
E1	9	59		E1	10	331

Table VIIC. HPV11 HLA-A1 Supermotif Peptides

E2	8	151
E2	9	151
E1	8	60
L2	8	152
El	11	436
Li	8	393
E5	10	45
Ll	8	344
L2	11	144
L2	9	38
E2	10	261
E2	8	354
E5	10	71
E5	11	71
L1	8	409
E1	10	207
E5	10	15
E5	9	50
E2	10	93
E6	8	26
L1	9 .	378
	10	389
L1 E6	10	11
	9	287
L2 L2		207
E2	10	149
	10	
E2	11	149 369
L2	11 9	151
L2		
E5	11 8	44 39
E6	10	39
E6		232
E1	11 9	585
E1		585
E1	11	37
L2	10	14
E5	11 10	150
L2 L2	11	149
		382
L2	11	
L1	8 9	224 332
E1		383
L2 L2	10	383
	11	78
E1	9	57
L1		
L1	11	57
E1	11	346 333
E1	8	
E1	11	499
E5	9	27
E5	9	32
E4	10	17
E1	8	275
L1	8	73
E5	11	48
E5	9	46
E2	11	128
L1	10	114
L1	9	62

f Peptid	les	
E2	9	206
L1	10	17
L1	8	297
L2	10	145
L2	11	414
E2	11	148
E1	10	584
L2	8	246
L1	9	72
E4	8	2
L1	8	58
L1	10	58
L2	9	120
E6	9	53
E5	10	14
E5	10	58
E5	11	58
E2	9	102
E2	8	92
E2	11	92
E4	8	22
L2	8	431
E1	9	579
E1	11	579
E2	8	138
L1	9	231
L1	8	246
L1	8	359
E5	10	61
E2	10	336
E2	9	159 159
E2	10	
L1	9	351
L1	8 9	26
L1		26
E4	11 8	16 9
E4 E2		168
E2 E1	11 8	502
E1	9	502
E2	8	131
	68091 v1	131
J. 11	00071 VI	

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotit-Bearing Peptides	
1 2	3	4	. HPV16 E1 8	548
HPV16 E1	9	316		10 548
HPV16 E1	11	316	HPV16 E1 8	3 75
HPV16 E1	9	239	HPV16 E1 9	9 75
HPV16 El	10	239	HPV16 E1 1	l1 75
HPV16 E1	8	317	HPV16 E1 8	3 22
HPV16 E1	10	317	HPV16 E1 1	11 22
HPV16 E1	10	205	HPV16 E1 9	374
HPV16 E1	8	478	HPV16 E1 1	0 374
HPV16 E1	11	478	HPV16 E1 9	356
HPV16 E1	10	112	HPV16 E1 1	10 213
HPV16 E1	11	112	HPV16 E1 1	11 213
HPV16 E1	9	539	HPV16 E1 8	65
HPV16 E1	11	539	HPV16 E1 9	65
HPV16 E1	8	69	HPV16 E1 8	63
HPV16 E1	9	459	HPV16 E1 1	LO 63
HPV16 E1	9	318	HPV16 E1 1	11 63
HPV16 E1	9	206	HPV16 E1 1	LO 288
HPV16 E1	10	73	HPV16 E1 1	1 288
HPV16 E1	11	73	HPV16 E1 8	140
HPV16 E1	10	380	HPV16 E1 8	3 138
HPV16 E1	10	406	HPV16 E1 1	LO 138
HPV16 E1	9	524	HPV16 E1 3	LO 331
HPV16 E1	10	82	HPV16 E1 9	51
HPV16 E1	11	82	HPV16 E1 1	LO 51
HPV16 E1	10	23		3 392
HPV16 E1	11	23	HPV16 E1 1	LO 392
HPV16 E1	11	405		11 392
HPV16 E1	8	237		11 463
HPV16 E1	11	237		LO 493
HPV16 E1	8	114		LO 445
HPV16 E1	9	114	HPV16 E1 9	
HPV16 E1	8	472	HPV16 E1 8	
HPV16 E1	10	472		LO 501
HPV16 E1	9	259	HPV16 E1 9	
HPV16 E1	10	259	HPV16 E1 8	
HPV16 E1	9	304	HPV16 E1 9 HPV16 E1 1	9 466 LO 466
HPV16 E1	8	187	HPV16 E1 B	
HPV16 E1	9	187		10 242
HPV16 E1	11	187	HPV16 E1 8	
HPV16 E1	8	353	HPV16 E1 8	
HPV16 E1	9	353 101	HPV16 E1 8	
HPV16 E1	10			571
HPV16 E1	9	640 640		LO 12
HPV16 E1	10 8	299	HPV16 E1	
HPV16 E1	9	299	HPV16 E1 8	
HPV16 E1 HPV16 E1	10	515	HPV16 E1 S	
HPV16 E1	11	515		10 450
HPV16 E1	10	523		11 450
HPV16 E1	11	81	HPV16 E1	
HPV16 E1	10	97		11 179
HPV16 E1	8	368		3 216
HPV16 E1	9	368		68
HPV16 E1	10	43		11 263
HPV16 E1	11	43		184
HPV16 E1	8	384		184
HPV16 E1	10	384		11 184
HPV16 E1	10	335	HPV16 E1	10 238
HPV16 E1	11	335	HPV16 E1	11 238
10 11				

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TABLE TAB Superinous South	ng r epinaes		
HPV16	E1	8	247	HPV16 E1	8	474
HPV16	E1	9	247	HPV16 E1	9	490
HPV16	E1	8	375	HPV16 E1	10	490
HPV16	E1	9	375	HPV16 E1	10	464
HPV16	El	11	375	HPV16 E1	11	464
HPV16		9	473	HPV16 E1	9	494
HPV16		10	194	HPV16 E1	9	346
HPV16		10	264	HPV16 E1	9	510
HPV16		11	264	HPV16 E1	11	510
HPV16		9	564	HPV16 E1	8	255
HPV16		8	369	HPV16 E1 HPV16 E1	10 9	255 145
HPV16		8	401	HPV16 E1	11	145
HPV16		10	442 52	HPV16 E1	8	457
HPV16		9	52	HPV16 E1	11	457
HPV16 HPV16		11	52	HPV16 E1	8	191
HPV16		11	204	HPV16 E1	10	191
HPV16		11	111	HPV16 E1	9	243
HPV16		8	517	HPV16 E1	11	243
HPV16		9	517	HPV16 E1	8	59
HPV16		10	517	HPV16 E1	9	59
HPV16		8	400	HPV16 E1	11	59
HPV16		9	400	HPV16 E1	9	554
HPV16	E1	8	296	HPV16 E1	10	554
HPV16	E1	10	296	HPV16 E1	11	554
HPV16	E1	11	296	HPV16 E1	11	222
HPV16	E1	9	292	HPV16 E1	11	544
HPV16		В	311	HPV16 E1	8	91
HPV16		9	311	HPV16 E1	10	306
HPV16		9	77	HPV16 E1	11	306
HPV16		10	77	HPV16 E1	8	207
HPV16		9	418	HPV16 E1	11	207 144
HPV16		10	117	HPV16 E1 HPV16 E1	10 8	305
HPV16		10	323	HPV16 E1	11	305
HPV16 HPV16		9 11	252 252	HPV16 E1	10	360
HPV16		8	199	HPV16 E1	11	360
HPV16	E1	9	199	HPV16 E1	11	569
HPV16	E1	10	199	HPV16 E1	8	202
HPV16		11	199	HPV16 E1	8	538
HPV16		8	267	HPV16 E1	10	538
HPV16		9	267	HPV16 E1	8	193
HPV16	E1	10	267	HPV16 E1	11	193
HPV16	E1	11	267	HPV16 E1	9	328
HPV16	E1	8	513	HPV16 E1	8	105
HPV16		9	513	HPV16 E1	9	105
HPV16		8	382	HPV16 E1	11	105
HPV16	E1	10	382	HPV16 E1	10	535
HPV16		10	208	HPV16 E1	11	535 136
HPV16	E1	8	563	HPV16 E1 HPV16 E1	9 10	136
HPV16		10	563	HPV16 E1	9	480
HPV16	E1	9	297	HPV16 E1	11	480
HPV16	E1	10	297 297	HPV16 E1	8	196
HPV16 HPV16		11 9	562	HPV16 E1	10	196
HPV16		11	562	HPV16 E1	11	196
HPV16	E1	9	254	HPV16 E1	10	4
HPV16	E1	11	254	HPV16 E1	9	512
HPV16		8	293	HPV16 E1	10	512
HPV16		11	293	HPV16 E1	8	561
TLATO						

Table VIII HLA-A2 Supermotif-Bearing Peptides

				oupermon bearing r	opia			
HPV16	E1	10	561	HP	V16	E1	10	458
HPV16	E1	9	94	HP	V16	E1	11	72
HPV16	E1	8	190		V16		8	185
HPV16	E1	9	190		V16		10	185
HPV16	E1	11	190		V16		11	185
HPV16	E1	10	553		V16		9.	289
HPV16		11	553				10	289
HPV16		11	302			E1	8	253
HPV16		11	636		V16		10	253
HPV16		9	61		V16		9	407
HPV16		10	61		V16		8 10	60 60
HPV16		9	398		V16 V16		11	60
HPV16		10	398			E1	11	344
HPV16		11	398 441		V16		8	525
HPV16 HPV16		9	381		V16		8	85
HPV16		11	381		V16		11	85
HPV16		8	556		V16		9	197
HPV16		9	556		V16		10	197
HPV16		10	556		V16		11	197
HPV16		11	143	HP	V16	E1	10	345
HPV16		8	419	HP	V16	E1	9	443
HPV16		11	359	HP	V16	E1	8	555
HPV16		9	256	HP	V16	E1	9	555
HPV16		8	188	HP	V16	E1	10	555
HPV16	E1	10	188		V16		11	555
HPV16	E1	11	188		V16		9 -	83
HPV16	E1	8	146		V16		10	83
HPV16		10	146		V16		9	361
HPV16		8	84		V16		10	361
HPV16		9	84			E1	9	24
HPV16		9	414		V16		10	24 363
HPV16		8	615		V16 V16		9	425
HPV16		11	432		V16		8	339
HPV16		10	390 246		V16		8	509
HPV16 HPV16		9	246		V16		10	509
HPV16		10	246		V16		11	379
HPV16		11	250		V16		9	531
HPV16		8	266	HP	V16	E1	10	531
HPV16		9	266	HP	V16	E1	8	261
HPV16		10	266	HP	V16	E1	8	578
HPV16		11	266	HP	V16	E1	10	578
HPV16	E1	8	484			E1	9	58
HPV16	E1	11	484		V16		10	58
HPV16	E1	10	489		V16		9	90
HPV16		11	489		V16		10	448
HPV16		8	634		V16		11	448
HPV16		9	546			E1	10	20
HPV16		10	546		V16		8	220
HPV16		10	397		V16		9 10	220
HPV16		11	397			E2	8	72
HPV16		11	423 314		V16	E2	10	72
HPV16		8	231		V16	E2	11	72
HPV16		9	231		V16	E2	9	41
HPV16 HPV16		10	231		V16		9	228
HPV16		10	315		V16		10	228
HPV16		8	66			E2	11	228
HPV16		11	66		V16	E2	9	69
11E V 20		-~						

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TIEA-A2 Supermont-Bears	ng r cpudes		
HPV16	E2	10	69	HPV16 E2	8	136
HPV16		11	69	HPV16 E2	10	136
HPV16		8	221	HPV16 E2	11	136
HPV16		9	221	HPV16 E2	8	214
HPV16		11	221	HPV16 E2	9	214
HPV16		9	226	HPV16 E2	11	214
HPV16		11	226	HPV16 E2	8	290
HPV16		8	63	HPV16 E2	9	290
HPV16		10	63	HPV16 E2	8	35
HPV16		11	63	HPV16 E2	8	56
HPV16		9	314	HPV16 E2	9	56
HPV16		10	40	HPV16 E2	9	223
HPV16		8	109	HPV16 E2	11	252
HPV16		9	109	HPV16 E2	11	210
HPV16		11	109	HPV16 E2	10	15
HPV16		11	300	HPV16 E2	10	238
				HPV16 E2	8	356
HPV16		11	5 309	HPV16 E2	10	356
HPV16		10		HPV16 E2	8	288
HPV16		10	174	HPV16 E2	10	288
HPV16		8	294	HPV16 E2	11	288
HPV16		9	124	HPV16 E2	8	68
HPV16		9	344	HPV16 E2	10	68
HPV16		8	246	HPV16 E2	11	68
HPV16		9	246	HPV16 E2	10	45
HPV16		11	246	HPV16 E2	10	225
HPV16		8	96			14
HPV16		9	96	HPV16 E2	11 8	351
HPV16		10	96	HPV16 E2 HPV16 E2	10	351
HPV16		11	142		8	255
HPV16		8	209	HPV16 E2 HPV16 E2	11	255
HPV16		8	74	HPV16 E2	10	354
HPV16		9	74	HPV16 E2	11	182
HPV16		11	4.8			215
HPV16		9	2	HPV16 E2 HPV16 E2	8 10	215
HPV16		8	185	HPV16 E2	8.	62
HPV16		9	185	HPV16 E2	9	62
HPV16		10	185	HPV16 E2	11	62
HPV16		8	118		10	256
HPV16		11	118	HPV16 E2 HPV16 E2	8	70
HPV16		8	204	HPV16 E2	9	70
HPV16		8	100	HPV16 E2	10	70
HPV16		11	100	HPV16 E2	8	94
HPV16		10	346	HPV16 E2	10	94
HPV16		11	346	HPV16 E2	11	94
HPV16		8	168	HPV16 E2	8	75
HPV16		9	156	HPV16 E2	8	103
HPV16		11	156	HPV16 E2	9	16
HPV16		8	150	HPV16 E2	11	16
HPV16		11	150		9	127
HPV16		8	190	HPV16 E2	11	127
HPV16		10	190	HPV16 E2	8	284
HPV16		8	230	HPV16 E2	8	284
HPV16		9	230	HPV16 E2		
HPV16		8	187	HPV16 E2	9	9
HPV16		11	187	HPV16 E2	8	325
HPV16		8	29	HPV16 E2	9	325
HPV16		10	29	HPV16 E2	10	325
HPV16		9	53	HPV16 E2	11	325
HPV16		10	53	HPV16 E2	8	219
HPV16	E2	11	53	HPV16 E2	9	219

Table VIII
HLA-A2 Supermotif-Bearing Peptides

HPV16	E2	10	219		HPV16	E2	8	227
HPV16	E2	11	219		HPV16		10	227
HPV16	E2	9	287		HPV16		11	227
HPV16	E2	11	287		HPV16	E2	8	192
HPV16	E2	11	106		HPV16	E2	10	119
HPV16	E2	10	60		HPV16	E2	11	119
HPV16		11	60		HPV16	E2	8	145
HPV16	E2	10	196		HPV16		11	147
HPV16		8	71			E2	10	341
HPV16	E2	9	71		HPV16	E2 E2	11	321 134
HPV16		11	71		HPV16		8	92
HPV16	E2	10	151		HPV16	E2 E2	10	92
HPV16	E2	9	191 .		HPV16	E2	8	138
HPV16		8	349		HPV16	E2	9	138
HPV16		9 10	349 349			E2	10	138
HPV16		8	57		HPV16		11	138
	E2	8	278		HPV16	E2	9	102
HPV16	E2	9	278			E2	11	312
	E2	11	278		HPV16	E2	9	131
HPV16		10	37		HPV16	E2	11	115
HPV16		9	7		HPV16	E2	8	159
HPV16		10	7		HPV16	E2	11	159
HPV16		11	7		HPV16	E5	9	53
HPV16		9	212		HPV16	E5	10	53
HPV16	E2	10	212		HPV16	E5	8	26
HPV16	E2	11	212		HPV16	E5	9	26
HPV16	E2	11	165		HPV16	E5	11	26
HPV16	E2	8	98		HPV16	E5	9	24
HPV16	E2	10	98		HPV16	E5	10	24
HPV16	E2	8	348		HPV16	E5	11	24
HPV16	E2	9	348		HPV16	E5	8	20
HPV16	E2	10	348		HPV16	E5	9	20
	E2	11	348		HPV16	E5	10	20
HPV16		9	85		HPV16	E5	8	5
HPV16		10	85		HPV16	E5	9	5 60
HPV16	E2	8	23		HPV16	E5 E5	8 10	60
HPV16	E2	10	317		HPV16	E5	10	72
HPV16		8	261		HPV16		11	72
HPV16	E2 E2	9 10	261 261		HPV16		8	15
HPV16		8	198		HPV16	E5	9	15
	E2	9	144		HPV16	E5	11	15
HPV16		11	269		HPV16	E5	8	66
	E2	10	313		HPV16		9	66
	E2	11	237		HPV16	E5	11	66
HPV16	E2	9	355		HPV16	E5	8	75
	E2	11	355		HPV16	E5	9	75
HPV16		9	61		HPV16	E5	8	64
HPV16		10	61		HPV16	E5	10	64
HPV16		8	3		HPV16	E5	11	64
HPV16	E2	9	93		HPV16	E5	9	43
HPV16	E2	11	93		HPV16	E5	10	43
HPV16	E2	9	310		HPV16	E5	11	43
HPV16	E2	8	128		HPV16	E5	8	44
HPV16	E2	10	128		HPV16	E5	9	44
HPV16	E2	10	253		HPV16	E5	10	44
HPV16	E2	11	285		HPV16	E5	11	44
	E2	10	116		HPV16		11	51
HPV16	E2	9	357		HPV16	E5	9	61

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				III.I III Gapen	 g - epilo			
HPV16	E5	11	61		HPV16	E5	8	9
HPV16	E5	10	12		HPV16	E5	9	9
HPV16	E5	11	12		HPV16	E5	8	21
HPV16	E5	9	73		HPV16	E5	9	21
HPV16	E5	10	73		HPV16	E5	8	46
HPV16	E5	11	73		HPV16	E5	9	46
HPV16	E5	ė.	42		HPV16	E5	11	46
HPV16	E5	10	42		HPV16	E5	9	63
HPV16	E5	11	42		HPV16	E5	11	63
HPV16	E5	9	28		HPV16	E5	9	68
HPV16	E5	11	28		HPV16	E5	11	68
HPV16	E5	11	11		HPV16		8	39
HPV16	E5	8	16		HPV16	E5	9	39
HPV16	E5	10	16		HPV16	E5	10	39
HPV16	E5	8	22		HPV16	E5	11	39
HPV16	E5	11	22		HPV16	E6	8	110
HPV16	E5	8	27		HPV16	E6	11	58
HPV16	E5	10	27		HPV16		8	73.
HPV16	E5	9	32		HPV16	E6	10	143
HPV16	E5	11	32		HPV16	E6	8	23
HPV16	E5	8	47		HPV16		11	23
HPV16	E5	10	47		HPV16		8	37
HPV16	E5	8	33		HPV16		9	37
HPV16		10	33		HPV16	E6	9	25
HPV16		11	33		HPV16		10	25
HPV16	E5	9	48		HPV16	E6	11	25
HPV16	E5	-8	45		HPV16	E6	8	96
HPV16		9	45		HPV16	E6	11	96
HPV16		10	45		HPV16	E6	10	48
HPV16		9	1		HPV16		8	52
HPV16		10	1		HPV16		9	52 9
HPV16		11	1		HPV16		11 11	125
HPV16		8	3		HPV16	E6		34
HPV16		9	3		HPV16		11 10	59
HPV16	E5	10	3		HPV16	E6	11	59
HPV16		11	3		HPV16		9	18
HPV16		9	70		HPV16		11	18
HPV16		8	31		HPV16		9	41
HPV16		10	31		HPV16		11	107
HPV16		8	55		HPV16		10	44
HPV16		10	55		HPV16		8	26
HPV16 HPV16		11	55 41			E6	9	26
		9	41		HPV16		10	26
HPV16		11	41		HPV16		11	134
HPV16 HPV16		9	8		HPV16		10	102
HPV16		10	8		HPV16	E6	11	116
HPV16		8	37		HPV16		8	12
HPV16		9	37		HPV16		11	12
HPV16		10	37		HPV16		9	20
HPV16		11	37		HPV16	E6	10	20
HPV16		8	35		HPV16	E6	11	20
HPV16		9	35		HPV16		8	21
HPV16		10	35		HPV16		9	21
HPV16		11	35		HPV16		10	21
HPV16		10	52		HPV16	E6	11	4.3
HPV16		11	52		HPV16	E6	8	42
HPV16		8	6		HPV16	E6	10	97
HPV16		11	6		HPV16	E6	11	97
HPV16		8	10		HPV16	E6	8	27
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Table VIII HLA-A2 Supermotif-Bearing Peptides

HPV16	E6	9	27	HPV16 E7	10	11
	E6	8	151	HPV16 L1	8	372
HPV16	E6	10	29	HPV16 L1	9	372
HPV16		10	94	HPV16 L1 HPV16 L1	8 11	451 451
HPV16		8	28	HPV16 L1	8	373
HPV16		11	28 93	HPV16 L1	9	233
HPV16		11	67	HPV16 L1	8	342
HPV16		9	68	HPV16 L1	10	330
	E7	11	68	HPV16 L1	8	513
HPV16		8	75	HPV16 L1	10	513
HPV16		9	75	HPV16 L1	11	513
	E7	10	75	HPV16 L1	8	35
	E7	9	81	HPV16 L1	10	35
HPV16		10	81	HPV16 L1	10	292
HPV16	E7	9	14	HPV16 L1	9	70
HPV16	E7	8	21	HPV16 L1	10	205
HPV16	E7	9	4	HPV16 L1	9	371
HPV16		10	4	HPV16 L1	10	371
HPV16	E7	9	37	HPV16 L1	9	172
HPV16		11	18	HPV16 L1	11	172
HPV16		8	43	HPV16 L1	9	183
HPV16		9	85	HPV16 L1	8	454
HPV16		10	73	HPV16 L1 HPV16 L1	11	251 329
HPV16		11	73	HPV16 L1	11 8	397
HPV16		11	54	HPV16 L1	11	397
HPV16 HPV16		9	82 82	HPV16 L1	10	300
HPV16		8	83	HPV16 L1	11	300
HPV16		11	83	HPV16 L1	9	225
HPV16		8	15	HPV16 L1	10	225
HPV16		8	12	HPV16 L1	10	486
HPV16		9	12	HPV16 L1	11	486
HPV16		11	12	HPV16 L1	9	154
HPV16	E7	10	41	HPV16 L1	10	154
HPV16	E7	8	6	HPV16 L1	10	228
HPV16	E7	10	6	HPV16 L1	11	228
HPV16	E7	11	44	HPV16 L1	8	120
HPV16		8	49	HPV16 L1	10	120
HPV16		9	66	HPV16 L1	9	113 361
HPV16		11	66	HPV16 L1 HPV16 L1	10	361
HPV16		10	77 77	HPV16 L1	10	442
HPV16		11	77	HPV16 L1	11	442
HPV16		8	71	HPV16 L1	9	412
HPV16		9	71	HPV16 L1	11	17
HPV16		10	63	HPV16 L1	9	34
HPV16		9	78	HPV16 L1	11	34
HPV16		10	78	HPV16 L1	8	279
HPV16		8	86	HPV16 L1	8	132
HPV16	E7	9	7	HPV16 L1	10	132
HPV16	E7	9	64	HPV16 L1	10	474
HPV16	E7	11	64	HPV16 L1	8	245
HPV16		10	19	HPV16 L1	10	245
HPV16		8	69	HPV16 L1	8	400
HPV16		10	69	HPV16 L1	9	400
	E7	11	69	HPV16 L1	10	400
	E7	10	55	HPV16 L1	11	400
	E7	11	55	HPV16 L1 HPV16 L1	9	5 494
HPV16	E7	9	11	MEATO PI	,	4.74

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-AZ Super	mont-Bearing Peptid	es		
HPV16	L1	8	402	HPV16	L1	10	406
HPV16	L1	9	402	HPV16	L1	8	151
HPV16		10	402	HPV16	L1	10	151
HPV16		11	25	HPV16	L1	11	262
HPV16		8	506	HPV16	L1	8	178
HPV16		9	506	HPV16	L1	9	90
HPV16		11	506	HPV16	L1	9	46
HPV16		11	236	HPV16	L1	10	46
HPV16	L1	9	282	HPV16	L1	9	312
HPV16		11	282	HPV16	L1	10	69
HPV16	L1	8	446	HPV16	L1	8	184
HPV16	L1	8	356	HPV16	L1	11	216
HPV16	L1	11	356	HPV16	L1	11	68
HPV16	L1	8	232	HPV16			148
HPV16	L1	10	232	HPV16			148
HPV16	L1	11	291	HPV16	L1	8	495
HPV16	L1	8	348	HPV16			239
HPV16	L1	10	348	HPV16			239
HPV16	L1	11	348	HPV16			398
HPV16	L1	8	142	HPV16			398
HPV16	L1	11	142	HPV16		8	432
HPV16	L1	9	499	HPV16		9	432
HPV16	L1	10	499	HPV16			339
HPV16	L1	9	431	HPV16			94
HPV16	L1	10	431	HPV16		10	94
HPV16		9	93	HPV16		9	409
HPV16		11	93	HPV16		10	9
HPV16		8	136	HPV16		8	87
HPV16		10	438	HPV16		8	124
HPV16		11	438	HPV16			124 1
HPV16		8	64	HPV16			1
HPV16		8	166	HPV16			1
HPV16		10	166 130	HPV16			1
HPV16		10 9	140	HPV16		10	414
HPV16 HPV16		10	140	HPV16		11	414
HPV16		8	62	HPV16		8	226
HPV16		9	62	HPV16			226
HPV16		10	62	HPV16			263
HPV16		8	22	HPV16		8	325
HPV16		10	22	HPV16		10	164
HPV16		8	285	HPV16	L1	9	157
HPV16		9	285	HPV16	L1	11	157
HPV16		11	457	HPV16	L1	8	58
HPV16		10	452	HPV16	L1	11	58
HPV16		9	424	HPV16	L1	10	311
HPV16	L1	11	8	HPV16	L1	8	476
HPV16	L1	9	86	HPV16		10	476
HPV16	L1	9	221	HPV16		8	367
HPV16	L1	8	11	HPV16		10	367
HPV16	L1	10	11	HPV16		8	353
HPV16		8	407	HPV16		10	353
HPV16	L1	9	407	HPV16		11	353
HPV16		11	407	HPV16		8	383
HPV16		8	501	HPV16		9	218
HPV16		9	512	HPV16		10	218
HPV16		11	512	HPV16		8	296
HPV16		10	85	HPV16		9	19
HPV16		8	406	HPV16 HPV16		11	19 460
HPV16	Ll	9	406	HLA19	n1	•	400

Table VIII HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bear	ing Peptides		
HPV16	T.1	10	77	HPV16 L1	9	365
HPV16		11	77	HPV16 L1	10	365
HPV16		8	247	HPV16 L1	10	375
HPV16		8	213	HPV16 L1	11	521
HPV16		9	213	HPV16 L1	8	410
HPV16		8	489	HPV16 L1	11	410
HPV16		10	489	HPV16 L1	9	523
HPV16		11	138	HPV16 L1	10	423
HPV16		10	466	HPV16 L1	9	439
HPV16		9	147	HPV16 L1	10	439
HPV16	L1	8	319	HPV16 L1	8	507
HPV16	L1	9	319	HPV16 L1	10	507
HPV16	L1	8	515	HPV16 L1	11	507
HPV16	L1	9	515	HPV16 L1	9	238
HPV16	L1	10	515	HPV16 L1	11	238
HPV16	L1	8	41	HPV16 L1	8	408
HPV16		10	41	HPV16 L1	10	408
HPV16		8	43	HPV16 L1	9	121
HPV16		11	497	HPV16 L1	11	121
HPV16		9	450	HPV16 L1	10	522
HPV16		9.	240	HPV16 L1	10	237
HPV16		11	240	HPV16 L1	9	362
HPV16		9	331	HPV16 L1	11	362
HPV16		8	403	HPV16 L1	8	516
HPV16		9	403	HPV16 L1	9	516
HPV16		11	403	HPV16 L1	9	219 219
HPV16		11	181	HPV16 L1 HPV16 L1	11	219
HPV16		9 .	354	HPV16 L1	9	358
HPV16		10	354 280	HPV16 L1	11	358
HPV16 HPV16		11	26	HPV16 L1	9	36
HPV16		11	26	HPV16 L1	10	54
HPV16		8	2	HPV16 L1	11	204
HPV16		9	2	HPV16 L1	8	220
HPV16		10	2	HPV16 L1	10	220
HPV16		11	2	HPV16 L1	9	10
HPV16		9	289	HPV16 L1	11	10
HPV16		9	341	HPV16 L1	8	413
HPV16		9	123	HPV16 L1	11	413
HPV16		11	123	HPV16 L1	8	3
HPV16	L1	8	56	HPV16 L1	9	3
HPV16	L1	10	56	HPV16 L1	10	3
HPV16	L1	9	482	HPV16 L1	10	357
HPV16	L1	9	159	HPV16 L1	8	359
HPV16		9	253	HPV16 L1	10	359
HPV16		11	253	HPV16 L1	8	47
HPV16		8	369	HPV16 L1	9	47
HPV16		11	369	HPV16 L1	11	47
HPV16		11	271	HPV16 L1	8	126
HPV16		8	28	HPV16 L1	8	30
HPV16		9	28	HPV16 L1	10	30
HPV16		10	28	HPV16 L1	8	416
HPV16		9	174	HPV16 L1	9	416
HPV16		11	174	HPV16 L1	10	416 302
HPV16		9	324	HPV16 L1 HPV16 L1	9	302
HPV16		10	449	HPV16 L1	11	302
HPV16		9	49	HPV16 L1	10	302
HPV16		11	49	HPV16 L1	11	38
HPV16		11	422 365	HPV16 L1	10	389
HPV16	T-T	8	303	HEATO DI	10	303

Table VIII HLA-A2 Supermotif-Bearing Peptides

			nt.A-A2 Superino	m-Bearing replices		
HPV16 L1	9	275		HPV16 L2	11	283
HPV16 L1	8	470		HPV16 L2	11	163
HPV16 L1	11	53		HPV16 L2	8	181
HPV16 L2	11	355		HPV16 L2	10	181
HPV16 L2	8	144		HPV16 L2	9	118
HPV16 L2	9	144		HPV16 L2	8	404
HPV16 L2	10	144		HPV16 L2	8	259
HPV16 L2	11	144		HPV16 L2	8	59
HPV16 L2	8	288		HPV16 L2	10	57
HPV16 L2	10	356		HPV16 L2	11	364
HPV16 L2	9	293		HPV16 L2	10	226
HPV16 L2	8	82		HPV16 L2	8	26
HPV16 L2	11	15		HPV16 L2	11	26
HPV16 L2	8	116		HPV16 L2	9	65
HPV16 L2	11	116		HPV16 L2	11	65
HPV16 L2	10	31		HPV16 L2	10	61
HPV16 L2	11	31		HPV16 L2	8	76
HPV16 L2	8	147		HPV16 L2	10	76
HPV16 L2	9	147		HPV16 L2	11	76
HPV16 L2	10	147		HPV16 L2	9	52
HPV16 L2	11	147		HPV16 L2	11	52
HPV16 L2	10	415		HPV16 L2 HPV16 L2	8 9	354 440
HPV16 L2	9	285		HPV16 L2	11	41
HPV16 L2	10	285		HPV16 L2	8	277
HPV16 L2	11 8	285 367		HPV16 L2	10	277
HPV16 L2 HPV16 L2	9	367		HPV16 L2	11	277
HPV16 L2	11	367		HPV16 L2	10	439
'HPV16 L2	9	422		HPV16 L2	9	32
HPV16 L2	10	422		HPV16 L2	10	32
HPV16 L2	9	43		HPV16 L2	11	32
HPV16 L2	11	43		HPV16 L2	8	145
HPV16 L2	11	199		HPV16 L2	9	145
HPV16 L2	10	84		HPV16 L2	10	145
HPV16 L2	11	84		HPV16 L2	11	145
HPV16 L2	10	376		HPV16 L2	9	4.5
HPV16 L2	9	140		HPV16 L2	8	420
HPV16 L2	8	129		HPV16 L2	9	420
HPV16 L2	9	129		HPV16 L2	11	420 374
HPV16 L2	11	129		HPV16 L2 HPV16 L2	9	344
HPV16 L2	8	338		HPV16 L2	9	344
HPV16 L2 HPV16 L2	11	338 195		HPV16 L2	8	243
HPV16 L2	9	195		HPV16 L2	9	243
HPV16 L2	11	195		HPV16 L2	8	135
HPV16 L2	9	340		HPV16 L2	10	135
HPV16 L2	11	340		HPV16 L2	11	135
HPV16 L2	8	176		HPV16 L2	11	250
HPV16 L2	9	111		HPV16 L2	8	286
HPV16 L2	11	111		HPV16 L2	9	286
HPV16 L2	8	114		HPV16 L2	10	286
HPV16 L2	10	114		HPV16 L2	9	430
HPV16 L2	8	373		HPV16 L2	10	430
HPV16 L2	10	373		HPV16 L2	11	430
HPV16 L2	8	242		HPV16 L2	8	105
HPV16 L2	9	242		HPV16 L2	11	105
HPV16 L2	10	242		HPV16 L2	8	202
HPV16 L2	9	201		HPV16 L2 HPV16 L2	9 10	202 202
HPV16 L2	10	201		HPV16 L2	9 .	248
HPV16 L2	11	201		MFV10 112	,	240

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TIER-NZ Supermont-Beath	15 replaces		
HPV16	L2	10	23	HPV16 L2	9	435
HPV16		11	23	HPV16 L2	8	80
HPV16		8	20	HPV16 L2	10	80
HPV16		8	39	HPV16 L2	8	161
HPV16		8	35	HPV16 L2	9	161
HPV16		11	35	HPV16 L2	11	246
HPV16		10	323	HPV16 L2	11	172
HPV16		11	323	HPV16 L2	8	358
HPV16		8	236	HPV16 L2	11	358
HPV16		9	236	HPV16 L2	11	120
HPV16		10	236	HPV16 L2	11	221
HPV16		8	86	HPV16 L2	9	97
HPV16		9	86	HPV16 L2	10	97
HPV16		10	86	HPV16 L2	8	381
HPV16		8	249	HPV16 L2	10	381
HPV16		9	169	HPV16 L2	8	88
HPV16		8	341	HPV16 L2	11	88
HPV16		10	341	HPV16 L2	9	24
HPV16		11	341	HPV16 L2	10	24
HPV16		8	46	HPV16 L2	8	423
HPV16		8	294	HPV16 L2	9	423
HPV16		8	108	HPV16 L2	8	44
HPV16		10	108	HPV16 L2	10	44
HPV16		9	410	HPV16 L2	9	17
HPV16		11	410	HPV16 L2	11	17
HPV16		9	454	HPV16 L2	9	233
HPV16		9	276	HPV16 L2	11	233
HPV16		11	276	HPV16 L2	9	342
HPV16		10	407	HPV16 L2	10	342
HPV16		9	419	HPV16 L2	11	342
HPV16		10	419	HPV16 L2	11	310
HPV16		9	397	HPV16 L2	8	234
HPV16		9	208	HPV16 L2	10	234
HPV16		8	150	HPV16 L2	11	234
HPV16		9	174	HPV16 L2	10	12
HPV16		10	174	HPV16 L2	8	305
HPV16		8	240	HPV16 L2	10	305
HPV16		10	240	HPV16 L2	8	224
HPV16		11	240	HPV16 L2	9	224
HPV16		9	143	HPV16 L2	9	461
HPV16		10	143	HPV16 L2	11	461
HPV16		11	143	HPV16 L2	9	298
HPV16		8	292	HPV16 L2	9	69
HPV16		10	292	HPV16 L2	8	9
HPV16		11	395	HPV16 L2	10	9
HPV16		8	255	HPV16 L2	8	313
HPV16		11	255	HPV16 L2	10	313
HPV16		8	417	HPV16 L2	8	230
HPV16		11	417	HPV16 L2	9	230
HPV16		8	215	HPV16 L2	9	335
HPV16		9	215	HPV16 L2	10	335
HPV16		11	215	HPV16 L2	11	335
HPV16		10	429	HPV16 L2	8	6
HPV16		11	429	HPV16 L2	10	6
HPV16		8	74	HPV16 L2	11	6
HPV16		9	74	HPV16 L2	8	14
HPV16		10	74	HPV16 L2	11	274
HPV16		8	409	HPV16 L2	9	360
HPV16		10	409	HPV16 L2	11	360
HPV16		9	197	HPV16 L2	11	125
**E A T O		-				

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Beari	ng Peptides		
HPV16	т э	8	134	HPV16 L2	9	351
HPV16		9	134	HPV16 L2	11	351
HPV16		11	134	HPV16 L2	9	136
HPV16		9	104	HPV16 L2	10	136
HPV16		8	389	HPV16 L2	11	136
HPV16		10	389	HPV16 L2	8	350
HPV16		11	389	HPV16 L2	10	350
HPV16		9	107	HPV16 L2	10	153
HPV16		11	107	HPV16 L2	8	209
HPV16		9	50	HPV16 L2	9	154
HPV16		11	50	HPV16 L2	11	154
HPV16	L2	8	138	HPV16 L2	8	287
HPV16	L2	9	138	HPV16 L2	9.	287
HPV16	L2	11	138	HPV16 L2	10	222
HPV16	L2	8	189	HPV16 L2	11	222
HPV16	L2	10	189	HPV16 L2	8	168
HPV16	L2	9	331	HPV16 L2	10	168
HPV16	L2	11	331	HPV16 L2	8	155
HPV16		11	186	HPV16 L2	10	155
HPV16		8	204	HPV16 L2	11	152
HPV16		11	204	HPV16 L2	8	237
HPV16		10	213	HPV16 L2	9	237
HPV16		11	213	HPV16 L2	11 9	237 369
HPV16		8	387	HPV16 L2 HPV16 L2	11	369
HPV16		10	387 378	HPV16 L2	8	393
HPV16 HPV16		8 11	378	HPV16 L2	10	72
HPV16		9	347	HPV16 L2	11	72
HPV16		10	347	HPV16 L2	8	447
HPV16		11	347	HPV16 L2	9	447
HPV16		9	167	HPV16 L2	10	453
HPV16		11	167	HPV16 L2	8	349
HPV16		9	122	HPV16 L2	9	349
HPV16	L2	11	384	HPV16 L2	11	349
HPV16	L2	9	81	HPV18 E1	11	396
HPV16	L2	8	332	HPV18 E1	10	397
HPV16		10	332	HPV18 E1	8	324
HPV16		11	438	HPV18 E1	10	324
HPV16		10	399	HPV18 E1	9	246 246
HPV16		10	187	HPV18 E1 HPV18 E1	10	246
HPV16		8	343 343	HPV18 E1	10	22
HPV16 HPV16		9 10	343	HPV18 E1	11	22
HPV16		9	85	HPV18 E1	9	546
HPV16		10	85	HPV18 E1	8	68
HPV16		11	85	HPV18 E1	9	466
HPV16		10	311	HPV18 E1	10	387
HPV16		9	182	HPV18 E1	11	387
HPV16		11	265	HPV18 E1	9	325
HPV16	L2	10	16	HPV18 E1	9	213
HPV16	L2	10	232	HPV18 E1	8	526
HPV16	L2	9	156	HPV18 E1	9	526
HPV16	L2	8	398	HPV18 E1	10	66
HPV16		11	398	HPV18 E1	8	72
HPV16		8	141	HPV18 E1	10	72
HPV16		11	141	HPV18 E1	11	72 422
HPV16		8	244	HPV18 E1 HPV18 E1	8 9	199
HPV16		10	2.5	HPV18 E1	8	40
HPV16 HPV16		8 11	231	HPV18 E1	9	40
45ATP	112	TT	231		-	

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TILATE TIL Dapormout Douring Copiesso
HPV18 E1	10	413	HPV18 E1 11 59
HPV18 E1	8	144	HPV18 E1 9 64
HPV18 E1	11	531	HPV18 E1 11 309
HPV18 E1	9	216	HPV18 E1 10 104
HPV18 E1	9	504	HPV18 E1 9 141
HPV18 E1	11	412	HPV18 E1 10 141
HPV18 E1	8	273	HPV18 E1 11 141
HPV18 E1	9	273	HPV18 E1 8 74
HPV18 E1	10	273	HPV18 E1 9 74
HPV18 E1	11	273	HPV18 E1 11 74
HPV18 E1	8	479	HPV18 E1 10 338
HPV18 E1	10	479	HPV18 E1 11 89
HPV18 E1	9	311	
HPV18 E1	10	404	
HPV18 E1	11	404	HPV18 E1 10 497 HPV18 E1 10 265
HPV18 E1	8	240	HPV18 E1 10 265
HPV18 E1	11	240	HPV18 E1 8 460
HPV18 E1	9	196	HPV18 E1 9 463
HPV18 E1	10	196	HPV18 E1
HPV18 E1	11	635	HPV18 E1 8 399
HPV18 E1	8	78	HPV18 E1 11 399
HPV18 E1	8	530 628	HPV18 E1 10 452
HPV18 E1		628	HPV18 E1 11 452
HPV18 E1	11 11	203	HPV18 E1 10 508
HPV18 E1 HPV18 E1	9	363	HPV18 E1 10 465
HPV18 E1	11	228	HPV18 E1 10 212
HPV18 E1	8	381	HPV18 E1 10 503
HPV18 E1	9	381	HPV18 E1 9 356
HPV18 E1	10	381	HPV18 E1 8 332
HPV18 E1	8	46	HPV18 E1 9 332
HPV18 E1	11	46	HPV18 E1 8 223
HPV18 E1	9	637	HPV18 E1 8 300
HPV18 E1	8	106	HPV18 E1 11 300
HPV18 E1	11	106	HPV18 E1 8 494
HPV18 E1	10	42	HPV18 E1 11 494
HPV18 E1	10	522	HPV18 E1 9 121
HPV18 E1	11	522	HPV18 E1 11 121
HPV18 E1	9	342	HPV18 E1 9 172
HPV18 E1	10	342	HPV18 E1 9 55
HPV18 E1	11	342	HPV18 E1 11 55
HPV18 E1	10	52	HPV18 E1 10 11
HPV18 E1	8	220	HPV18 E1 8 473
HPV18 E1	10	220	HPV18 E1 9 473
HPV18 E1	11	220	HPV18 E1 8 182
HPV18 E1	8	540	HPV18 E1 11 182 HPV18 E1 8 279
HPV18 E1	11	30	
HPV18 E1	8	166	
HPV18 E1	8	143	
HPV18 E1	9	143	HPV18 E1 11 270 HPV18 E1 8 83
HPV18 E1	11	115	HPV18 E1 8 83 HPV18 E1 8 306
HPV18 E1	8	62	HPV18 E1 9 306
HPV18 E1	11	62	HPV18 E1 8 254
HPV18 E1	9	108	HPV18 E1 8 254 HPV18 E1 9 254
HPV18 E1	11	108	HPV18 E1 8 198
HPV18 E1	8	375	HPV18 E1 10 198
HPV18 E1	9	375	HPV18 E1 10 1569
HPV18 E1	11	366	HPV18 E1 9 266
HPV18 E1	8	59	HPV18 E1 10 271
HPV18 E1	10	59	HEVIO 21 10 2/1

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-	Bearing Peptid	es		
HPV18	F1	11	271	HPV18	E1	9	520
HPV18		9	501		E1	8	350
HPV18		B	562	HPV18	E1	8	571
HPV18		9	562	HPV18	E1	9	295
HPV18		10	562		E1	10	295
HPV18		11	562	HPV18	E1	11	295
HPV18	E1	8	262	HPV18	E1	9	480
HPV18		10	262	HPV18	E1	10	229
HPV18		10 .	314	HPV18	E1	11	229
HPV18		11	314	HPV18	E1	8	382
HPV18		11	347	HPV18	E1	9	382
HPV18		11	461	HPV18	E1	8	214
HPV18		9	590 -	HPV18	E1	11	214
HPV18		9	23	HPV18	E1	8	527
HPV18	E1	10	23	HPV18	E1	11	527
HPV18	E1	10	449	HPV18	E1	8	312
HPV18		8	124	HPV18	E1	10	47
HPV18	E1	9	124 -	HPV18	E1	10	367
HPV18	E1	11	439	HPV18	E1	11	367
HPV18	E1	11	647	HPV18	E1	8	545
HPV18	E1	8	318	HPV18	E1	10	545
HPV18	E1	9	318	HPV18	E1	9	39
HPV18	E1	8	210	HPV18	E1	10	39
HPV18	E1	8	259	HPV18	E1	10	188
HPV18	E1	9	259	HPV18	E1	11	188
HPV18	E1	11	259	HPV18		9	335
HPV18		8	237	HPV18	E1	9	487
HPV18		9	237	HPV18	E1	8	158
HPV18		10	237	HPV18	E1	10	158
		11	237	HPV18	E1	11	158
HPV18	E1	8	524	HPV18 HPV18	E1 E1	9	191 191
HPV18		9	524 524	HPV18	E1	11	191
HPV18 HPV18	E1	10	524	HPV18	E1	10	577
HPV18		8	206	HPV18	E1	11	485
HPV18	E1	9	206	HPV18	E1	8	568
HPV18	E1	10	206		E1	11	568
HPV18		11	206	HPV18		11	551
HPV18		8	389		E1	11	448
	E1	9	389	HPV18	E1	8	98
		10	389	HPV18	E1	10	98
	E1	10	215	HPV18	E1	10	560
HPV18	E1	9	561	HPV18	E1	11	560
HPV18	E1	10	561	HPV18	E1	8	519
HPV18	E1	11	561	HPV18	E1	9	519
HPV18	E1	9	261	HPV18	E1	10	519
HPV18	E1	11	261	HPV18		8	194
HPV18	E1	11	313	HPV18		11	194
HPV18		9	388	HPV18		9	252
	E1	10	388	HPV18		10	252
HPV18	E1	11	388	HPV18		11	252
HPV18		9	304	HPV18		9	60
HPV18		10	304		E1	10	60
HPV18		11	304	HPV18		8	21
HPV18		10	204	HPV18 HPV18	E1	11	21 405
	E1	11	204		E1	10	405
HPV18		11 9	285 570		E1	11	405
	E1 E1	8	376		E1	9	67
HPV18		8	520	HPV18		8	457
11E A T 9		-				-	

Table VIII HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bearing	ng Peptide	:S		
HPV18	E1	10	457	HPV18	E1.	10	150
HPV18		11	457	HPV18		10	532
HPV18		8	563	HPV18		8	296
HPV18		9	563	HPV18		9	296
HPV18		10	563	HPV18		10	296
HPV18		11	563	HPV18		8	591
HPV18		8	200	HPV18		9	323
HPV18		8	426	HPV18	E1	11	323
HPV18		9	456	HPV18		8	297
HPV18		11	456	HPV18		9	297
HPV18		11	80	HPV18		11	297
HPV18		9	649	HPV18		8	525
HPV18		9	421	HPV18		9	525
HPV18		10	589	HPV18	E1	10	525
HPV18		11	626	HPV18		10	31
HPV18		8	102	HPV18		11	31
HPV18		9	349	HPV18		8	505
HPV18		8	294	HPV18		10	81
HPV18		10	294	HPV18		11	84
HPV18		11	294	HPV18	E1	9	339
HPV18		9	425	HPV18	E1	9	20
HPV18		10	330	HPV18	E1	9	450
HPV18		11	330	HPV18		9	368
HPV18		8	622	HPV18		10	368
HPV18		9	553	HPV18		10	244
HPV18		10	553	HPV18	E1	11	244
HPV18		9	117	HPV18	E1	11	149
HPV18		11	430	HPV18	E1	8	370
HPV18		10	164	HPV18		8	346
HPV18		11	93	HPV18	E1	9	432
HPV18		9	302	HPV18	E1	8	516
HPV18		11	302	HPV18	E1	10	516
HPV18		10	511	HPV18	E1	11	516
HPV18		11	511	HPV18	E1	8	536
HPV18		10	322	HPV18	E1	11	536
HPV18		11	179	HPV18	E1	8	243
HPV18		9	245	HPV18	E1	11	243
HPV18		10	245	HPV18	E1	11	386
HPV18	E1	11	245	HPV18	E1	8	585
HPV18		8	65	HPV18	E1	8	408
HPV18		11	65	HPV18	E1	11	542
HPV18	E1	8	253	HPV18	E1	8	455
HPV18	E1	9	253	HPV18	E1	10	455
HPV18	E1	10	253	HPV18	E1	10	19
HPV18	E1	8	197	HPV18	E2	9	49
HPV18	E1	9	197	HPV18	E2	10	49
HPV18	E1	11	197	HPV18	E2	10	245
HPV18	E1	8	260	HPV18	E2	11	245
HPV18	E1	10	260	HPV18	E2	8	76
HPV18	E1	8	303	HPV18	E2	11	76
HPV18	E1	10	303	HPV18	E2	11	45
HPV18	E1	11	303	HPV18		8	351
HPV18	E1	9	414	HPV18		9	351
HPV18	E1	9	53	HPV18		10	351
HPV18	E1	11	53	HPV18		11	87
HPV18	E1	8	238	HPV18		9	154
HPV18	E1	9	238	HPV18		8	214
HPV18	E1	10	238	HPV18		11	214
HPV18	E1	9	533	HPV18		9	246
HPV18	E1	11	533	HPV18	E2	10	246

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Beam	ng Peptides		
HPV18	E2	10	132	HPV18 E2	8	329
HPV18		10	156	HPV18 E2	8	238
HPV18		8	146	HPV18 E2	9	238
HPV18		9	146	HPV18 E2	10	238
HPV18	E2	10	146	HPV18 E2	10	254
HPV18		11	29	HPV18 E2	8	86
HPV18		9	315	HPV18 E2	8	39
HPV18		11	315	HPV18 E2	11	39
HPV18	E2	9	100	HPV18 E2	8	266
HPV18	E2	8	210	HPV18 E2	8	98
HPV18	E2	9	210	HPV18 E2	11	98
HPV18	E2	9	78	HPV18 E2	11	83
HPV18		10	78	HPV18 E2	11	221
HPV18	Ē2	8	104	HPV18 E2	8	79
HPV18		10	6	HPV18 E2	9	79
HPV18		8	340	HPV18 E2	9	333
HPV18		10	340	HPV18 E2	8	217
HPV18		11	340	HPV18 E2	8	1 .
HPV18		8	190	HPV18 E2	10	144
HPV18		9	190	HPV18 E2	11 9	144
HPV18		8	48	HPV18 E2 HPV18 E2	10	67
HPV18		10	4.8	HPV18 E2	11	67
HPV18		11	48	HPV18 E2	8	285
HPV18		11	346	HPV18 E2	9	348
HPV18		9	324	HPV18 E2	11	348
HPV18		10 11	324 324	HPV18 E2	9	196
HPV18 HPV18		11	331	HPV18 E2	10	64
HPV18		9	54	HPV18 E2-	9	265
HPV18		10	54	HPV18 E2	10	272
HPV18		11	253	HPV18 E2	11	110
HPV18		9	85	HPV18 E2	8	262
HPV18		11	161	HPV18 E2	9	262
HPV18		9	235	HPV18 E2	10	262
HPV18		11	235	HPV18 E2	8	357
HPV18		8	148	HPV18 E2	9	357
HPV18		10	148	HPV18 E2	8	33
HPV18		11	187	HPV18 E2	8	38
HPV18	E2	9	291	HPV18 E2	9	38
HPV18	E2	9	60	HPV18 E2	9	216
HPV18	E2	9	223	HPV18 E2	8	80
HPV18	E2	10	223	HPV18 E2	8	56
HPV18	E2	11	289	HPV18 E2	10	56
HPV18	E2	10	332	HPV18 E2	11	2
HPV18		8	358	HPV18 E2	8	61
HPV18		8	55	HPV18 E2	9	11
HPV18		9	55	HPV18 E2	10 8	11 343
HPV18		11	55	HPV18 E2 HPV18 E2	9	343
HPV18		8	72	HPV18 E2	10	343
HPV18		10	72		11	244
HPV18		11	72	HPV18 E2 HPV18 E2	9	213
HPV18		8	75	HPV18 E2	9	298
HPV18		9	75	HPV18 E2	9	203
HPV18		8	280	HPV18 E2	10	203
HPV18		10	280 280	HPV18 E2	8	32
HPV18		11	257	HPV18 E2	9	32
HPV18		10 11	257	HPV18 E2	9	206
HPV18 HPV18		11	152	HPV18 E2	10	206
HPV18		10	92	HPV18 E2	8	230
U5AT8	54	10				

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				ILA-Az Sup	ermoth-beath	g i cpuu	us .		
HPV18	E2	10	230			HPV18	E5	8	57
HPV18		8	318			HPV18	E5	10	57
HPV18	E2	11	233			HPV18	E5	9 .	50
HPV18	E2	9	355			HPV18	E5	10	50
HPV18		10	355			HPV18	E5	8	37
HPV18		11	355			HPV18	E5	11	37
HPV18	E2	8	140			HPV18	E5	8	65
HPV18		10	140			HPV18	E5	8	19
HPV18		8	236			HPV18	E5	10	19
HPV18		10	236			HPV18	E5	9	43
HPV18		11	236			HPV18	E5	10	43
HPV18		10	153			HPV18	E5	8	40
HPV18		9	57			HPV18	E5	9	40
HPV18		9	97.			HPV18	E5	10	40
HPV18		9	7			HPV18	E5	9	4
HPV18		10	215			HPV18	E5	11	4
HPV18		9	341			HPV18	E5	8	63
HPV18		10	341			HPV18	E5	10	63
HPV18		11	341			HPV18	E5	8	62
HPV18	E2	8	349			HPV18	E5	9	62
HPV18	E2	10	349			HPV18	E5	11	62
HPV18		11	349			HPV18	E5	9	58
HPV18		8	211			HPV18	E5	11	58
HPV18	E2	11	211			HPV18	E5	9	22
HPV18		9	231			HPV18	E5	11	22
HPV18		8	334			HPV18	E5	8	35
HPV18		11	334			HPV18	E5	9	35
HPV18	E2	9	350			HPV18	E5	10	35
HPV18		10	350		-	HPV18	E5	8	61
HPV18	E2	11	350			HPV18	E5	9	61
HPV18		9	136			HPV18	E5	10	61
HPV18	E2	10	136			HPV18	E5	9	46
HPV18	E2	8	197			HPV18		8	21
HPV18	E2	11	197			HPV18	E5	10	21
HPV18	E2	8	356			HPV18		9	60
HPV18	E2	9	356			HPV18		10	60
HPV18	E2	10	356			HPV18		11	60
HPV18	E2	10	335			HPV18		10	3
HPV18	E2	9	37			HPV18		8	25
HPV18	E2	10	37			HPV18		10	25
HPV18	E2	9	322			HPV18		11	25
HPV18	E2	11	322			HPV18		11	48
HPV18		10	96			HPV18		8	51
HPV18		11	143			HPV18		9	51
HPV18		10	135			HPV18		11	51
HPV18		11	135			HPV18		8	42
HPV18		8	164			HPV18		10	42
HPV18		11	164			HPV18		11	42
HPV18		8	47			HPV18		9	34
HPV18		8	29			HPV18		10	34
HPV18		10	29			HPV18			34
HPV18		8	27			HPV18 HPV18		11	41
HPV18		9	27					9	41
HPV18		10	27			HPV18 HPV18		11	41
HPV18		8	13			HPV18		8	33
HPV18		10	13			HPV18		9	33
HPV18		11	13			HPV18		10	33
HPV18		10	11			HPV18		11	33
HPV18		9	6			HPV18		8	31
HPV18	E5	11	6			UBATS	22		31

Table VIII
HLA-A2 Supermotif-Bearing Peptides

HPV18 E5	10	31	HPV18 E6 9	95
HPV18 ES	11	31	HPV18 E6 9	22
HPV18 E5	9	39	HPV18 E6 10	22
HPV18 E5	10	39	HPV18 E6 8	114
HPV18 E5	11	39	HPV18 E6 8	111
HPV18 E5	8	15	HPV18 E6 11	111
HPV18 E5	9	15	HPV18 E6 8	7
HPV18 E5	9	53	HPV18 E6 11	7
HPV18 E5	10	53	HPV18 E6 8	149
HPV18 E5	11	53	HPV18 E6 10	149
HPV18 E6	8	68	HPV18 E6 11	146
HPV18 E6	11	68	HPV18 E6 11	59
HPV18 E6	8	105	HPV18 E6 8	24 24
HPV18 E6	11	105	HPV18 E6 10	84
HPV18 E6	8	108	HPV18 E6 10 HPV18 E6 11	84
HPV18 E6	11	108	HPV18 E6 8	89
HPV18 E6	8	18	HPV18 E6 8 HPV18 E6 10	89
HPV18 E6	11	18	HPV18 E6 8	37
HPV18 E6	8	32	HPV18 E6 0	38
HPV18 E6	10	32	HPV18 E6 10	54
HPV18 E6	11	32 27	HPV18 E6 11	54
HPV18 E6 HPV18 E6	11 8	16	HPV18 E7 8	6
HPV18 E6	10	16	HPV18 E7 10	6
HPV18 E6	10	51	HPV18 E7 8	63
HPV18 E6	9	88	HPV18 E7 10	63
HPV18 E6	11	88	HPV18 E7 8	24
HPV18 E6	9	29	HPV18 E7 8	82
HPV18 E6	10	29	HPV18 E7 10	82
HPV18 E6	11	29	HPV18 E7 8	69
HPV18 E6	9	20	HPV18 E7 10	40
HPV18 E6	11	20	HPV18 E7 8	90
HPV18 E6	9	77	HPV18 E7 8	86
HPV18 E6	9	40	HPV18 E7 9	86
HPV18 E6	10	43	HPV18 E7 9	43
HPV18 E6	8	47	HPV18 E7 8	14
HPV18 E6	9	47	HPV18 E7 10	14 46
HPV18 E6	8	53	HPV18 E7 9 HPV18 E7 11	11
HPV18 E6	11	53	HPV18 E7 8	5
HPV18 E6	10	97 136	HPV18 E7 9	5
HPV18 E6	10	62	HPV18 E7 11	5
HPV18 E6 HPV18 E6	8 11	120	HPV18 E7 8	73
HPV18 E6	8	30	HPV18 E7 11	73
HPV18 E6	9	30	HPV18 E7 8	8
HPV18 E6	10	30	HPV18 E7 10	74
HPV18 E6	9	13	HPV18 E7 10	61
HPV18 E6	11	13	HPV18 E7 11	92
HPV18 E6	10	92	HPV18 E7 11	50
HPV18 E6	11	92	HPV18 E7 9	17
HPV18 E6	9	36	HPV18 E7 10	17
HPV18 E6	11	102	HPV18 E7 9	56
HPV18 E6	9	25	HPV18 E7 10	22
HPV18 E6	9	150	HPV18 E7 10	88
HPV18 E6	8	41	HPV18 E7 8	87
HPV18 E6	9	93	HPV18 E7 11	87 53
HPV18 E6	10	93	. HPV18 E7 8 HPV18 E7 9	53
HPV18 E6	11	93	HPV18 E7 9 HPV18 E7 10	53
HPV18 E6	8	1	HPV18 E7 10 HPV18 E7 8	84
HPV18 E6	8	95	HPVIO L/	3.

Table VIII
HLA-A2 Supermotif-Bearing Pentides

			HLA-A2 Supermotif-Bearing	ng Peptides		
HPV18	D-2	10	84	HPV18 L1	8	167
	E7	11	84	HPV18 L1	10	167
	E7	10	71	HPV18 L1	8	155
	E7	11	79	HPV18 L1	10	155
HPV18		9	7	HPV18 L1	10	280
HPV18		10	93	HPV18 L1	9	317
HPV18		11	60	HPV18 L1	11	317
HPV18		10	12	HPV18 L1	8	436
HPV18		9	75	HPV18 L1	9	436
HPV18		11	75	HPV18 L1	10	436
HPV18		10	195	HPV18 L1	11	436
HPV18		10	225	HPV18 L1	10	49
HPV18		11	225	HPV18 L1	8	438
HPV18		8	487	HPV18 L1	9	438
HPV18		9	487	HPV18 L1	10	438
HPV18		11	487	HPV18 L1	8	482
HPV18		9	63	HPV18 L1	8	391
HPV18		10	63	HPV18 L1	11	391
HPV18		10	268	HPV18 L1	8	267
HPV18		8	377	HPV18 L1	11	267
HPV18		11	419	HPV18 L1	8	535
HPV18	L1	9	196	HPV18 L1	11	177
HPV18	L1	8	552	HPV18 L1	10	342
HPV18	L1	11	552	HPV18 L1	8	171
HPV18	L1	10	222	HPV18 L1	9	233
HPV18	L1	8	406	HPV18 L1	11	326
HPV18	L1	8	218	HPV18 L1	8	383
HPV18	L1	9	218	HPV18 L1	10	383
HPV18		9	310	HPV18 L1	11	383
HPV18		8	2	HPV18 L1	10	165 467
HPV18		9	2	HPV18 L1	8 10	467
HPV18		8	490	HPV18 L1 HPV18 L1	11	467
HPV18		11	286	HPV18 L1	11	194
HPV18		9	441	HPV18 L1	8	97
HPV18		10	441 350	HPV18 L1	9	97
HPV18 HPV18		11	512	HPV18 L1	10	97
HPV18		10	512	HPV18 L1	9	38
HPV18		8	433	HPV18 L1	10	38
HPV18		10	433	HPV18 L1	11	38
HPV18		11	433	HPV18 L1	9	13
HPV18		9	260	HPV18 L1	10	428
HPV18		10	260	HPV18 L1	8	40
HPV18	L1	10	522	HPV18 L1	9	40
HPV18	L1	11	522	HPV18 L1	11	40
HPV18	L1	8	189	HPV18 L1	8	39
HPV18	L1	9	189	HPV18 L1	9	39
HPV18		11	263	HPV18 L1	10	39
HPV18	L1	8	276	HPV18 L1	8	46
HPV18		10	276	HPV18 L1	10	46
HPV18		9	148	HPV18 L1	9	460
HPV18		8	396	HPV18 L1	9	47 219
HPV18		9	396	HPV18 L1	9	9
HPV18		10	396	HPV18 L1 HPV18 L1	8	32
HPV18		9	330	HPV18 L1	9	32
HPV18		9	478 478	HPV18 L1	10	32
HPV18		10 11	478	HPV18 L1	8	488
HPV18		9	4/8	HPV18 L1	10	488
HPV18		8	203	HPV18 L1		443
UBATR	11		200			

Table VIII HLA-A2 Supermotif-Bearing Peptides

				-0F		
HPV18	L1.	11	443	HPV18 L1	8	282
HPV18	L1	8	360	HPV18 L1	8	248
HPV18	L1	9	376	HPV18 L1	9	248
HPV18	L1	10	186	HPV18 L1	8	525
HPV18	L1	11	186	HPV18 L1	10	525
HPV18		9	505	HPV18 L1	9 .	28
HPV18	L1	9	120	HPV18 L1	10	28
HPV18	L1	8	213	HPV18 L1	8	26
HPV18	L1	11	213	HPV18 L1	9	26 .
HPV18		9	125	HPV18 L1	11	26
HPV18	L1	8	8	HPV18 L1	10	240
HPV18	L1	10	8	HPV18 L1	8	20
HPV18		8	14	HPV18 L1	10	20
HPV18	Ll	11	103	HPV18 L1	9	333
HPV18		8	274	HPV18 L1	11	333
HPV18	L1	10	274	HPV18 L1	10	540
HPV18	L1	9	434	HPV18 L1	8	91
HPV18		10	434	HPV18 L1	8	472
HPV18		11	434	HPV18 L1	8	412
HPV18	L1	9	445	HPV18 L1	9	412
HPV18	L1	11	403	HPV18 L1	9	533
HPV18		10	104	HPV18 L1	10	533
HPV18		8	476	HPV18 L1	8	216
HPV18		11	476	HPV18 L1	10	216
HPV18	L1	11	531	HPV18 L1	11	216
HPV18	L1	8	159	HPV18 L1	8	439
HPV18	L1	10	159	HPV18 L1	9	439
HPV18		8	33	HPV18 L1	11	439
HPV18		9	33	HPV18 L1	11	315
HPV18		10	62	HPV18 L1	9	366
HPV18	L1	11	62	HPV18 L1	9	389
HPV18		8	261	HPV18 L1	10	389
HPV18		9	261	HPV18 L1	10	137
HPV18		11	36	HPV18 L1	11	61
HPV18		8	402	HPV18 L1	11	297
HPV18		8	388	HPV18 L1	10	214
HPV18		10	388	HPV18 L1	8	324
HPV18		11	388	HPV18 L1	9	324
HPV18		9	84	HPV18 L1	9	158
HPV18		11	84	HPV18 L1	11	158
HPV18		9	253	HPV18 L1	9	6
HPV18		10	253	HPV18 L1	10	6
HPV18		10	70	HPV18 L1	9	81
HPV18		11	70	HPV18 L1	10	81 299
HPV18		10	510	HPV18 L1	9	
HPV18		9	54	HPV18 L1	9 10	551 127
HPV18		10	54	HPV18 L1	9	288
HPV18		11	54	HPV18 L1 HPV18 L1	11	288
HPV18		9	52	HPV18 L1	11	93
HPV18		11	52	HPV18 L1	10	459
HPV18		10	199			31
HPV18		9	207	HPV18 L1 HPV18 L1	9 10	31
HPV18		11	207	HPV18 L1		31
HPV18		11	496	HPV18 L1	11 9	359
HPV18		10	114	HPV18 L1	10	150
HPV18		8	224	HPV18 L1	11	150
HPV18		11	224	HPV18 L1	9	518
HPV18		9	558	HPV18 L1	9	475
HPV18		8	344	HPV18 L1	9	335
HPV18	L1	8	57	ULATO DI	,	222

Table VIII
HI.A-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bearing	ng Peptides		
HPV18	T.1	11	335	HPV18 L1	10	394
HPV18		11	306	HPV18 L1	11	394
HPV18			242	HPV18 L1	В	82
HPV18		10	242	HPV18 L1	9	82
HPV18		10	365	HPV18 L1	11	82
HPV18			272	HPV18 L1	8	161
HPV18			400	HPV18 L1	9	452
HPV18		10	400	HPV18 L1	10	452
HPV18		10	485	HPV18 L1	9	45
HPV18		11	485	HPV18 L1	11	45
HPV18		8	78	HPV18 L1	9	337
HPV18		9	209	HPV18 L1	8	73
HPV18		11	209	HPV18 L1	10 .	73
HPV18		8	234	HPV18 L1	11	73
HPV18		8	446	HPV18 L1	8	129
HPV18		11	446	HPV18 L1	10	129
HPV18		10	404	HPV18 L1	11	4
HPV18		9	541	HPV18 L1	11	88
HPV18		8	442	HPV18 L2	9	6
HPV18		9	442	HPV18 L2	10	6 '
HPV18		9	273	HPV18 L2	8	286
HPV18		11	273	HPV18 L2	9	286
HPV18		10	444	HPV18 L2	10	341
HPV18		10	327	HPV18 L2	9	303
HPV18		9	215	HPV18 L2	11	303
HPV18		11	215	HPV18 L2	10	139
HPV18		9	156	HPV18 L2	9	358
HPV18		11	156	HPV18 L2	10	358
HPV18		11	409	HPV18 L2	9	278
HPV18		8	397	HPV18 L2	10	278
HPV18		9	397	HPV18 L2	11	278
HPV18		11	397	HPV18 L2	8	404
HPV18		11	473	HPV18 L2	10	404
HPV18		10	553	HPV18 L2	9	142
HPV18		9	105	HPV18 L2	11	142
HPV18		11	105	HPV18 L2	8	129
HPV18		8	254	HPV18 L2	9	129
HPV18		9	254	HPV18 L2	11	129
HPV18		11	254	HPV18 L2	10	349
HPV18		8	331	HPV18 L2	11	349
HPV18		11	331	HPV18 L2	10	346
HPV18		9	393	HPV18 L2	11	16
HPV18	L1	11	393	HPV18 L2	8	354
HPV18	L1	9	71	HPV18 L2	9	83
HPV18	L1	10	71	HPV18 L2	10	83
HPV18	L1	9	486	HPV18 L2	11	83
HPV18	L1	10	486	HPV18 L2	8	270
HPV18	L1	11	79	HPV18 L2	10	270
HPV18	L1	8	255	HPV18 L2	11	270
HPV18	L1	10	255	HPV18 L2	10	396
HPV18	L1	8	7	HPV18 L2	11	396
HPV18	L1	9	7	HPV18 L2	9	30
HPV18	L1	11	7	HPV18 L2	10	30
HPV18		8	449	HPV18 L2	11	30
HPV18		10	532	HPV18 L2	8	194
HPV18	L1	11	532	HPV18 L2	8	334
HPV18	L1	11	136	HPV18 L2	9	334
HPV18	L1	10	89	HPV18 L2	8	175
HPV18	L1	10	3 9 2	HPV18 L2	10	175
HPV18	L1	8	3 94	HPV18 L2	8	169

Table VIII HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Beari	ng Peptides		
HPV18	т э	9	169	HPV18 L	2 8	416
HPV18		8	455	HPV18 L		416
HPV18		9	369	HPV18 L		103
HPV18		10	369	HPV18 L		103
HPV18		11	200	HPV18 L		43
HPV18		9	443	HPV18 L		43
HPV18		9	53	HPV18 L		22
HPV18		8	241	HPV18 L		22
HPV18		9	241	HPV18 L	2 8	19
HPV18		10	241	HPV18 L	2 8	34
HPV18		11	276	HPV18 L	2 11	34
HPV18		9	122	HPV18 L	2 11	40
HPV18	L2	10	122	HPV18 L	2 8	106
HPV18	L2	11	157	HPV18 L	29	106
HPV18	L2	8	306	HPV18 L	2 8	248
HPV18	L2	10	306	HPV18 L		335
HPV18	L2	9	181	HPV18 L		197
HPV18	L2	8	116	HPV18 L		45
HPV18	L2	10	116	HPV18 L		45
HPV18		10	314	HPV18 L		263
HPV18		9	51	HPV18 L		242
HPV18		11	51	HPV18 L		242 287
HPV18		8	58	HPV18 L		391
HPV18		9	429	HPV18 L		391
HPV18		10	56	HPV18 L		338
HPV18		8	300	HPV18 L		79
HPV18		8 11	25 25	HPV18 L		179
HPV18		10	204	HPV18 L		179
HPV18		9	64	HPV18 L		254
HPV18		11	64	HPV18 L		254
HPV18		10	60	HPV18 L	2 10	254
HPV18		8	188	HPV18 L	2 11	254
HPV18		10	188	HPV18 L	2 8	160
HPV18		9	432	HPV18 L	29	160
HPV18	L2	8	310	HPV18 L	2 11	160
HPV18	L2	8	124	HPV18 L		285
HPV18	L2	10	124	HPV18 L		285
HPV18	L2	8	37	HPV18 L		422
HPV18		9	37	HPV18 L		138
HPV18	L2	8	134	HPV18 L		357
HPV18		10	134	HPV18 L		357
HPV18		11	134	HPV18 L		325 325
HPV18		8	292	HPV18 L:		209
HPV18		8	326	HPV18 L		209
HPV18		10	326	HPV18 L		415
HPV18		10	167 167	HPV18 L		415
HPV18		11	279	HPV18 L		73
HPV18		9	279	HPV18 L		73
HPV18		10	279	HPV18 L		73
HPV18		9	44	HPV18 L		214
HPV18		11	44	HPV18 L		214
HPV18		9	405	HPV18 L		196
HPV18		8	143	HPV18 L	2 8	390
HPV18		10	143	HPV18 L	2 10	390
HPV18		8	130	HPV18 L		337
HPV18		10	130	HPV18 L		171
HPV18		11	130	HPV18 L		419
HPV18	L2	11	249	HPV18 L	2 8	98

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bearing	ng Peptides		
HPV18	1.2	9	98	HPV18 L2	9	409
HPV18		10	98	HPV18 L2	8	235
HPV18		11	120	HPV18 L2	9	235
HPV18		8	86	HPV18 L2	10	149
HPV18		11	86	HPV18 L2	8	13
			185	HPV18 L2	11	111
HPV18		11	216	HPV18 L2	9	412
HPV18		11 9	95	HPV18 L2	10	412
HPV18			95	HPV18 L2	9	420
HPV18		10	95	HPV18 L2	11	420
HPV18		11	360	HPV18 L2	11	377
HPV18		8	360	HPV18 L2	8	105
HPV18		11	90 .	HPV18 L2	9	105
HPV18		10	398	HPV18 L2	10	105
HPV18		9	398	HPV18 L2	8	406
HPV18		10	398	HPV18 L2	11	406
HPV18		11	232	HPV18 L2	10	262
HPV18		8	198	HPV18 L2	8	304
		9	172	HPV18 L2	10	304
HPV18		11	172	HPV18 L2	9	425
HPV18			233	HPV18 L2	8	38
HPV18		10		HPV18 L2	11	261
HPV18		11	233 5	HPV18 L2	8	154
HPV18		8	5	HPV18 L2	8	136
HPV18		10	5	HPV18 L2	9	136
HPV18		11	11	HPV18 L2	8	410
HPV18		10	302	HPV18 L2	11	410
HPV18		10	229	HPV18 L2	9	135
HPV18		9	229	HPV18 L2	10	135
HPV18		8	298	HPV18 L2	10	388
HPV18 HPV18		10	298	HPV18 L2	11	293
HPV18		8	281	HPV18 L2	10	217
HPV18		10	225	HPV18 L2	8	80
HPV18		11	220	HPV18 L2	9	176
HPV18		8	316	HPV18 L2	11	176
HPV18		11	316	HPV18 L2	10	221
HPV18		11	450	HPV18 L2	8	236
HPV18		8	132	HPV18 L2	8	92
HPV18		9	132	HPV18 L2	9	140
HPV18		10	132	HPV18 L2	11	140
HPV18		8	380	HPV18 L2	9	104
HPV18		9	380	HPV18 L2	10	104
HPV18		10	380	HPV18 L2	11	104
HPV18		8	340	HPV18 L2	9	113
HPV18		11	340	HPV18 L2	11	113
HPV18		8	166	HPV18 L2	11	387
HPV18		11	166	HPV18 L2	11	81
HPV18		8	151	HPV18 L2	9	91
HPV18		11	151	HPV18 L2	8	31
HPV18		11	102	HPV18 L2	9	31
HPV18		9	49	HPV18 L2	10	31
HPV18		11	49	HPV18 L2	11	31
HPV18		9	247	HPV18 L2	10	112
HPV18		10	212	HPV18 L2	8	351
HPV18		11	212	HPV18 L2	9	351
HPV18		10	424	HPV18 L2	11	351
HPV18		8	147	HPV18 L2	8	332
HPV18		9	147	HPV18 L2	10	332
HPV18		9	153	HPV18 L2	11	332
HPV18		8	409	HPV18 L2	11	427
		-				

Table VIII HLA-A2 Supermotif-Bearing Peptides

			HLA-A2	Supermotif-Bearn	ng Peptid	es		
HPV18	T.2	10	71		HPV31	E1	8	528
HPV18		11	71		HPV31		10	528
HPV18		9	436		HPV31		8	348
HPV18		8	400		HPV31		9	348
HPV18		11	400		HPV31		10	311
HPV31		9	296		HPV31		8	74
HPV31		11	296		HPV31		9	74
HPV31		8	219		HPV31		11	74
HPV31		9	219		HPV31		8	62
HPV31		10	219		HPV31		11	62
HPV31		8	297		HPV31		8	21
HPV31		10	297		HPV31		11	80
HPV31		.10	185		HPV31		9	354
HPV31		10	111		HPV31	E1	10	354
HPV31		11	111		HPV31		10	127
HPV31		9	519		HPV31		8	193
HPV31		11	519		HPV31		10	193
HPV31		8	68		HPV31		11	193
HPV31		9	439		HPV31		9	64
HPV31		10	533		HPV31	E1	10	315
HPV31		11	533		HPV31	E1	11	315
HPV31		9	298		HPV31		8	168
HPV31	E1	9	186		HPV31	E1	10	168
HPV31		10	66		HPV31	E1	11	168
HPV31		8	72		HPV31	E1	8	139
HPV31		10	72		HPV31	E1	8	137
HPV31		11	72		HPV31	E1	10	137
HPV31	E1	10	360		HPV31	E1	11	443
HPV31	E1	9	504		HPV31	E1	8	372
HPV31	E1	11	22		HPV31	E1	10	372
HPV31	E1	10	81		HPV31	E1	11	372
HPV31	E1	10	370		HPV31	E1	10	473
HPV31	E1	8	263		HPV31		10	425
HPV31	E1	11	263		HPV31		9	436
HPV31	E1	8	113		HPV31		9	206
HPV31	E1	9	113		HPV31		8	433
HPV31	E1	10	452		HPV31		10	433
HPV31		8	279		HPV31		8	499
HPV31	E1	9	279		HPV31		8	467
HPV31		9	239		HPV31		8	305
HPV31		10	239		HPV31		8	252
HPV31		9	284		HPV31		11	403
HPV31		8	213		HPV31		10	11
HPV31		11	213		HPV31		10	160 386
HPV31		8	217		HPV31 HPV31		10 9	225
HPV31		10	217		HPV31		10	225
HPV31		11	217		HPV31		11	225
HPV31		10	100 620		HPV31		8	446
HPV31		-	620		HPV31		9	446
HPV31 HPV31		10 10	495		HPV31		10	446
			495		HPV31		8	196
HPV31 HPV31		11 10	503		HPV31		8	78
HPV31		10	96		HPV31		9	71
HPV31		11	421		HPV31		11	71
HPV31		9	336		HPV31		11	243
HPV31		11	46		HPV31		8	355
HPV31		10	42		HPV31		9	355
HPV31		9	332		HPV31		9	453
HPV31		10	332		HPV31		9	287
Pv31		10				-	-	

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TILA-A2 Supe	mon-bearing reput	cs		
HPV31	E1	10	287	HPV31	E1	11	175
HPV31		11	287	HPV31	E1	11	258
HPV31	E1	10	268	HPV31	E1	8	187
HPV31	E1	11	268	HPV31	E1	11	187
HPV31	E1	8	381	HPV31		8	285
HPV31	E1	10	422	HPV31	E1	11	285
HPV31	E1	11	184	HPV31		10	255
HPV31	E1	11	110	HPV31		8	257
HPV31	E1	11	532	HPV31		8	535
HPV31	E1	8	497	HPV31		9	535
HPV31	El	9	497	HPV31		10	535
HPV31		10	497	HPV3 1		11	535
HPV31		8	380	HPV31		10	47
HPV31		9	380	HPV31		9	143
HPV31		10	276	HPV31 HPV31		10 11	143
HPV31		11	276	HPV31		10	340
HPV31		9	272	HPV31		11	340
HPV31		11	272 291	HPV31		11	549
HPV31		9	291	HPV31		8	518
HPV31 HPV31		10	119	HPV31		10	518
		9	232	HPV31		8	173
HPV31 HPV31		8	179	HPV31		9	308
HPV31		9	179	HPV31		8	104
HPV31		10	179	HPV31		9	104
HPV31		11	412	HPV31		11	104
HPV31		8	247	HPV31		8	59
HPV31		9	247	HPV31	E1	10	59
HPV31		10	247	HPV31	E1	11	59
HPV31		11	247	HPV31	E1	9	135
HPV31	E1	8	493	HPV31	E1	10	135
HPV31	E1	9	493	HPV31	E1	9	460
HPV31	E1	8	362	HPV31		11	460
HPV31	E1	10	362	HPV31		9	55
HPV31	E1	8	454	HPV31		11	55
HPV31		10	286	HPV31		9	4
HPV31		11	286	HPV31		9	492
HPV31		11	202	HPV31		10	492
HPV31		9	470	HPV31		8	541 541
HPV31		10	470	HPV31 HPV31		10 9	93
HPV31		8	543	HPV31		8	170
HPV31		9	277	HPV31		9	170
HPV31		10	277 277	HPV31		11	170
HPV31 HPV31		11	273	HPV31		10	524
HPV31		10	273	HPV31		11	524
HPV31		9	542	HPV31		9	60
HPV31		11	234	HPV31		10	60
HPV31		9	256	HPV31		10	378
HPV31		9	534	HPV31	E1	11	378
HPV31		10	534	HPV31	E1	9	67
HPV31		11	534	HPV31	El	8	430
HPV31		9	474	HPV31	E1	10	430
HPV31		8	326	HPV31		11	430
HPV31		9	326	HPV31		9	361
HPV31	E1	9	490	HPV31		11	361
HPV31	E1	11	490	HPV31		8	536
HPV31	E1	10	235	HPV31		9	536
HPV31		10	244	HPV31		10	536
HPV31	E1	11	244	HPV31	El	8	399

Table VIII HLA-A2 Supermotif-Bearing Peptides

				 -67			
HPV31	E1	10	142	HPV31	E1	10	177
HPV31	E1	11	142	HPV31	E1	11	177
HPV31	E1	11	339	HPV31	E1	9	325
HPV31	E1	9	429	HPV31	E1	10	325
HPV31	E1	11	429	HPV31	E1	8	349
HPV31	E1	11	141	HPV31	E1	11	254
HPV31	E1	11	323	HPV31	E1	8	144
HPV31	E1	8	145	HPV31	E1	9	144
HPV31	E1	9	145	HPV31	E1	10	144
HPV31	E1	8	83	HPV31	E1	9	82
HPV31	E1	10	176	HPV31	E1	9	341
HPV31	E1	11	176	HPV31		10	341
HPV31	E1	9	394	HPV31		11	223
HPV31	E1	8	267	HPV31	E1	8	343
HPV31	E1	11	267	HPV31		8	319
HPV31	E1	9	398	HPV31	E1	9	405
HPV31	E1	10	303	HPV31		8	489
HPV31	E1	8	595	HPV31		10	489
HPV31	E1	10	438	HPV31		10	481
HPV31	E1	8	526	HPV31		11	359
HPV31	E1	9	526	HPV31		9	511
HPV31	E1	10	526	HPV31		10	511
HPV31	E1	8	246	HPV31		8	558
HPV31		9	246	HPV31		10	558
HPV31		10	246	HPV31		11	515
HPV31		11	246	HPV31		10	428
HPV31		10	469	HPV31		10	19
HPV31		11	469	HPV31		9	89 277
HPV31		11	377	HPV31 HPV31		8 10	277
HPV31		11	294	HPV31		11	277
HPV31		8	211	HPV31		9	278
HPV31		9 10	211 211	HPV31		10	278
HPV31		11	616	HPV31		8	72
HPV31 HPV31		10	295	HPV31		10	72
HPV31		9	120	HPV31		11	72
HPV31		8	65	HPV31		8	338
HPV31		11	65	HPV31		10	338
HPV31		9	269	HPV31		8	229
HPV31		10	269	HPV31		11	229
HPV31		8	233	HPV31	E2	9	69
HPV31		10	152	HPV31	E2	10	69
HPV31		9	387	HPV31	E2	11	69
HPV31	E1	8	333	HPV31	E2	9	61
HPV31	E1	9	333	HPV31	E2	10	61
HPV31	E1	11	151	HPV31		8	291
HPV31	E1	8	505	HPV31		10	239
HPV31	E1	8	226	HPV31		8	286
HPV31	E1	9	226	HPV31		10	286
HPV31	E1	10	226	HPV31		11	286
HPV31	E1	10	324	HPV31		9	228
HPV31	E1	11	324	HPV31		8	140
HPV31	E1	9	218	HPV31		9	140
HPV31		10	218	HPV31		8	109
HPV31		11	218	HPV31		9	109
HPV31		8	227	HPV31		11	109
HPV31		9	227	HPV31		9 10	33 0 330
HPV31		10	23	HPV31		11	330
HPV31		11	84	HPV31		8	280
HPV31	E1	9	177	HPV31	E 2	0	200

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TIEA-AZ Supermont-Bearin	ig i epitues		
HPV31	E2	8	145	HPV31 E2	11	210
HPV31		10	40	HPV31 E2	9	339
HPV31		8	301	HPV31 E2	8	66
HPV31		9	124	HPV31 E2	10	66
HPV31		8	204	HPV31 E2	8	68
HPV31		9	204	HPV31 E2	10	68
HPV31		11	204	HPV31 E2	11	68
HPV31		8	74	HPV31 E2	10	45
HPV31		9	74	HPV31 E2	8	358
HPV31		8	100	HPV31 E2	10	358
HPV31		11	100	HPV31 E2	8	260
HPV31		11	48	HPV31 E2	11	260
HPV31		10	320 .	HPV31 E2	8	213
HPV31		9	2	HPV31 E2	9	213
HPV31		8	185	HPV31 E2	10	213
HPV31		9	185	HPV31 E2	10	316
HPV31		10	185	HPV31 E2	11	226
HPV31		8	118	HPV31 E2	10	261
HPV31		11	118	HPV31 E2	8	42
HPV31		8	207	HPV31 E2	10	42
HPV31		10	207	HPV31 E2	8	70
HPV31		11	207	HPV31 E2	9	70
HPV31		11	136	HPV31 E2	10	70
HPV31		10	353	HPV31 E2	8	75
HPV31		11	353	HPV31 E2	11	75
HPV31		10	171	HPV31 E2	8	103
HPV31			168	HPV31 E2	8	78
		9	50	HPV31 E2	9	77
HPV31 HPV31		10	50	HPV31 E2	8	94
HPV31		8	209	HPV31 E2	10	94
		9	209	HPV31 E2	11	94
HPV31		10	156	HPV31 E2	9	337
HPV31 HPV31		11	156	HPV31 E2	11	337
		10	143	HPV31 E2	10	303
HPV31			190	HPV31 E2	10	282
HPV31		10 8		HPV31 E2	11	282
HPV31 HPV31		11	150	HPV31 E2	10	84
HPV31		8	179	HPV31 E2	11	84
		10	179	HPV31 E2	8	254
HPV31 HPV31		9	231	HPV31 E2	9	254
HPV31		10	231	HPV31 E2	11	127
HPV31		11	231	HPV31 E2	9	219
HPV31		9	273	HPV31 E2	11	219
HPV31		9	235	HPV31 E2	8	355
HPV31		8	187	HPV31 E2	9	355
HPV31		8	29	HPV31 E2	11	355
HPV31		10	29	HPV31 E2	10	361
HPV31		8	35	HPV31 E2	11	361
HPV31		9	35	HPV31 E2	8	9
HPV31		9	164	HPV31 E2	8	60
HPV31		8	297	HPV31 E2	10	60
HPV31		9	297	HPV31 E2	11	60
HPV31		9	56	HPV31 E2	9	290
HPV31		8	295	HPV31 E2	9	294
HPV31		10	295	HPV31 E2	11	294
		11	295	HPV31 E2	8	215
HPV31		9	304	HPV31 E2	11	106
HPV31 HPV31		8	165	HPV31 E2	8	71
HPV31		11	165	HPV31 E2	9	71
HPV31		8	210	HPV31 E2	11	71
nPV31	EZ.	o	210	/		

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			nLA-A2 Supermont-Bearn	ig repudes		
HPV31	E2	9	317	HPV31 E2	11	220
HPV31		10	76	HPV31 E2	10	116
HPV31		9	95	HPV31 E2	8	356
HPV31		10	95	HPV31 E2	10	356
HPV31		11	95	HPV31 E2	9	362
HPV31		9	283	HPV31 E2	10	362
HPV31		10	283	HPV31 E2	11	362
HPV31		11	283	HPV31 E2	8	274
HPV31	E2	8	96	HPV31 E2	11	274
HPV31		9	96	HPV31 E2	8	192
HPV31	E2	10	96	HPV31 E2	9	41
HPV31	E2	9	191	HPV31 E2	11	41
HPV31	E2	10	151	HPV31 E2	10	119
HPV31	E2	9	321	HPV31 E2	11	119
HPV31	E2	11	321	HPV31 E2	10	211
HPV31	E2	8	57	HPV31 E2	11	211
HPV31	E2	11	57	HPV31 E2	8	340
HPV31	E2	11	238	HPV31 E2	11	147
HPV31	E2	8	285	HPV31 E2	10	58
HPV31	E2	9	285	HPV31 E2	11	328
HPV31	E2	11	285	HPV31 E2	8	92
HPV31		10	37	HPV31 E2	10	92
HPV31	E2	9	7	HPV31 E2	11	344
HPV31	E2	10	7	HPV31 E2	9	138
HPV31		8	311	HPV31 E2	10	138
HPV31		9	247	HPV31 E2	11	138
HPV31		9	276	HPV31 E2	9	102
HPV31		11	276	HPV31 E2	9	131
HPV31		9	53	HPV31 E2	11	131
HPV31		10	53	HPV31 E2	11	115
HPV31		8	98	HPV31 E2	8	159
HPV31		10	98	HPV31 E2	11	159 40
HPV31		9	348	HPV31 E5	9	40
HPV31		10	348	HPV31 E5 HPV31 E5	10	40
HPV31		11	5	HPV31 E5	10	53
HPV31		9 11	346 346	HPV31 E5	9	61
HPV31		8	324	HPV31 E5	11	61
HPV31		9	266	HPV31 E5	8	26
HPV31 HPV31		10	266	HPV31 E5	9	26
HPV31		8	198	HPV31 E5	11	26
HPV31		11	198	HPV31 E5	8	20
HPV31		9	269	HPV31 E5	9	20
HPV31		10	269	HPV31 E5	10	20
HPV31		11	269	HPV31 E5	9	3
HPV31		8	63	HPV31 E5	10	3
HPV31		10	63	HPV31 E5	11	3
HPV31		11	63	HPV31 E5	8	66
HPV31		8	364	HPV31 ES	9	66
HPV31		9	364	HPV31 E5	11	66
HPV31		8	3	HPV31 E5	8	15
HPV31		10	128	HPV31 E5	9	15
HPV31		9	93	HPV31 E5	11	15
HPV31		11	93	HPV31 E5	9	24
HPV31		11	292	HPV31 E5	10	24
HPV31		9	221	HPV31 E5	11	24
HPV31	E2	10	221	HPV31 E5	10	72
HPV31	E2	9	240	HPV31 E5	11	52
HPV31	E2	8	220	HPV31 E5	10	48
HPV31	E2	10	220	HPV31 E5	8	46

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TILA-AZ Supermont-Dearm	ig i cpudes		
HPV31	E5	9	46	HPV31 E5	8	33
HPV31	E5	11	11	HPV31 E5		33
HPV31	E5	8	45	HPV31 E5		33
HPV31	E5	9	45	HPV31 E5		33
HPV31	E5	10	45	HPV31 E5		64
HPV31	E5	8	16	HPV31 E5		64
HPV31	E5		16	HPV31 E5		64
HPV31			22	HPV31 E5	-	50
HPV31			22	HPV31 E5		39
HPV31			44	HPV31 E5		39
HPV31			44	HPV31 E5		39
HPV31			44	HPV31 E5		39
HPV31			44	HPV31 E5 HPV31 E5		68 68
HPV31			43	HPV31 E5		63
HPV31			43 43	HPV31 E5	11	63
HPV31			42	HPV31 B6		18
HPV31 HPV31			42 -	HPV31 E6	11	18
HPV31			42	HPV31 E6		136
HPV31			27	HPV31 E6	8	103
HPV31			27	HPV31 E6		66
HPV31			32	HPV31 E6	11	63
HPV31			32	HPV31 E6	8	30
HPV31			32	HPV31 E6	9	30
HPV31		8	1	HPV31 E6	11	30
HPV31	E5	9	1	HPV31 E6	8	98
HPV31	E5	11	1	HPV31 E6		49
HPV31	E5		5	HPV31 E6	8	57
HPV31			5	HPV31 E6-	11	57
HPV31			70	HPV31 E6	9	20
HPV31			56	HPV31 E6	8	14
HPV31			56	HPV31 E6	8 10	39 39
HPV31			31	HPV31 E6 HPV31 E6	8	41
HPV31			31	HPV31 E6	10	41
HPV31 HPV31		11 8	31 10	HPV31 E6	11	41
HPV31			7	HPV31 E6	8	45
HPV31			7	HPV31 E6	9	45
HPV31		8	35	HPV31 E6	10	95
HPV31		9	35	HPV31 E6	11	95
HPV31		10	35	HPV31 E6	8	35
HPV31		11	35	HPV31 E6	9	35
HPV31		8	37	HPV31 E6	8	85
HPV31	E5	9	37	HPV31 E6	11	118
HPV31	E5	10	37	HPV31 E6	9	137
HPV31	E5	11	37	HPV31 E6	11	137
HPV31		8	41	HPV31 E6	11	52
HPV31		9	41	HPV31 E6	8	11
HPV31		11	41	HPV31 E6	9	11
HPV31		9	8	HPV31 E6	11	11
HPV31		10	8	HPV31 E6	10	90
HPV31		9	73	HPV31 E6 HPV31 E6	11 11	90 100
HPV31		8	47	HPV31 E6	10	37
HPV31		11	47	HPV31 E6	9	50
HPV31		9	28 28	HPV31 E6	9	91
HPV31		11	28 12	HPV31 E6	10	91
HPV31		11	12	HPV31 E6	11	91
HPV31 HPV31		8	21	HPV31 E6	11	127
HPV31		9	21	HPV31 E6	8	5
nrv31	E-3	,	41		-	-

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			The composition board	-6 r spinass		
HPV31	E6	11	5	HPV31 E7	11	72
HPV31	E6	11	109	HPV31 E7	9	37
HPV31	E6	8	36	HPV31 E7	8	12
HPV31	E6	11	36	HPV31 E7	9	12
HPV31	E6	11	27	HPV31 E7	11	12
HPV31	E6	10	17	HPV31 E7	8	69
HPV31		10	82	HPV31 E7	10	69
HPV31	E6	11	82	HPV31 E7	11	69
HPV31	E6	8	51	HPV31 E7	10	55
HPV31	E6	10	87	HPV31 E7	11	55
HPV31	E6	11	86	HPV31 E7	9	11
HPV31		9	42	HPV31 E7	10	11
HPV31		10	42	HPV31 L1	8	347
HPV31		11	42	HPV31 L1	9	347
HPV31		10	19	HPV31 L1	8	348
HPV31		9	59	HPV31 L1	10	398
HPV31		11	59	HPV31 L1	11	398
HPV31		9	68 -	HPV31 L1	8	426
HPV31		11	68	HPV31 L1	10	180
HPV31		8	75	HPV31 L1	9	213
HPV31		9	75	HPV31 L1	11	213
HPV31		10	75	HPV31 L1	10	208
HPV31		8	21	HPV31 L1	8	317
HPV31		9	14	HPV31 L1	10	305
HPV31		8	48	HPV31 L1	11	285
HPV31		9	48	HPV31 L1 HPV31 L1	8 10	9
HPV31		10	36	HPV31 L1		346
HPV31		11	18	HPV31 L1	9 10	346
HPV31		9	81 81	HPV31 L1	9	147
HPV31		10 9	4	HPV31 L1	11	147
HPV31 HPV31		10	4	HPV31 L1	9	158
HPV31		9	88	HPV31 L1	11	304
HPV31		11	88	HPV31 L1	9	387
HPV31		8	89	HPV31 L1	8	372
HPV31		10	89	HPV31 L1	11	372
HPV31		11	54	HPV31 L1	10	275
HPV31		8	82	HPV31 L1	11	275
HPV31		9	82	HPV31 L1	9	200
HPV31		8	83	HPV31 L1	10	200
HPV31		8	8	HPV31 L1	10	461
HPV31		11	79	HPV31 L1	11	461
HPV31		8	15	HPV31 L1	9	129
HPV31		9	41	HPV31 L1	10	203
HPV31		8	6	HPV31 L1	11	203
HPV31	E7	10	6	HPV31 L1	8	216
HPV31	E7	11	44	HPV31 L1	10	216
HPV31	E7	11	27	HPV31 L1	9	88
HPV31	E7	10	73	HPV31 L1	8	336
HPV31	E7	11	73	HPV31 L1	10	336
HPV31	E7	8	77	HPV31 L1	10	417
HPV31	E7	9	66	HPV31 L1	11	417
HPV31	E7	11	66	HPV31 L1	9	8
HPV31	E7	10	63	HPV31 L1	11	8
HPV31	E7	8	71	HPV31 L1	8	95
HPV31	E7	9	71	HPV31 L1	10	95
HPV31	E7	9	7	HPV31 L1	8	107
HPV31	E7	9	64	HPV31 L1	10	107
HPV31		11	64 .	HPV31 L1	10	449
HPV31	E7	8	72	HPV31 L1	8	375

Table VIII HLA-A2 Supermotif-Bearing Peptides

HPV31 L1	9 10 8 10 11	373 373 69 69 407
HPV31 L1	8 10 9 10 8 10 11	69 69 407 43
HPV31 L1	10 9 10 8 10 11	69 407 43
HPV31 L1	9 10 8 10 11	407 43
HPV31 L1	10 8 10 11 10	43
HPV31 L1	8 10 11 10	
HPV31 L1	10 11 10	
HeV31 L1	11 10	99
HPV31 L1	10	99
HPV31 L1		314
HPV31 L1		389 389
HPV31 L1		238
HPV31 L1		238
HPV31 L1 1 323		201
HOV31 L1		201
HPV31 L1		300
HPV31 L1		179
HPV31 L1	9	32
HPV31 L1	11	32
HPV31 L1	8	451
HPV31 L1	10	451
HPV31 L1	8 `	342
HPV31 L1		342
		328
HPV31 L1		328
HPV31 L1		328
HPV31 L1		220
HPV31 L1		397
HPV31 L1 9 399		222 188
HPV31 L1		188
HPV31 L1		464
HPV31 L1 R 388		464
HPV31 L1		113
		122
HPV31 L1		294
HPV31 L1 9 382	9	294
HPV31 L1	. 8	15
IPV31 L1 11 181 IPV31 L1 IPV31 L1 IPV31 L1 IPV31 L1 10 482 IPV31 L1 IPV31 L1	10	15
HPV31 L1 9 61 HPV31 L1		17
HPV31 L1		425
HPV31 L1 8 381 HPV31 L1 HPV31 L1 HPV31 L1 HPV31 L1 1 10 381 HPV31 L1 HPV31 L1 HPV31 L1 HPV31 L1 10 60 HPV31 L1 HPV31 L1 10 237 HPV31 L1 HPV31 L1 11 237 HPV31 L1 HPV31 L1 11 237 HPV31 L1		425
HPV31 L1 9 381		306
HPV31 L1 10 381 HPV31 L1 HPV31 L1 HPV31 L1 10 60 HPV31 L1 10 237 HPV31 L1 HPV31 L1 11 237 HPV31 L1		378
HPV31 L1 10 60 HPV31 L1 HPV31 L1 10 237 HPV31 L1 HPV31 L1 11 237 HPV31 L1		378 156
HPV31 L1 10 237 HPV31 L1 HPV31 L1 1237 HPV31 L1		329
HPV31 L1 11 237 HPV31 L1		329
11.131 31 11		316
HPV31 L1 8 153 HPV31 L1		476
HPV31 L1 8 153 HPV31 L1 HPV31 L1 HPV31 L1		98
HPV31 L1 9 20 HPV31 L1		98
HPV31 L1 10 20 HPV31 L1		30
HPV31 L1 9 287 HPV31 L1		30
HPV31 L1 8 159 HPV31 L1		385
HPV31 L1 8 123 HPV31 L1	. 11	385
HPV31 L1 8 470 HPV31 L1	. 9	457
HPV31 L1 11 42 HPV31 L1	. 8	487
		487
HPV31 L1 8 214 HPV31 L1 HPV31 L1 HPV31 L1		487

Table VIII
HLA-A2 Supermotif-Bearing Peptides

HPV31	L1	8	490		V31 L1	11	21
HPV31	L1	9	228		V31 L1	8	101
HPV31	L1	11	228		V31 L1	8 .	391
HPV31		11	51		V31 L1	9	391
HPV31	L1	9	414		V31 L1	10	391
HPV31	Ll	10	414		V31 L1	8	277
HPV31		8	2		V31 L1	9	277
HPV31	L1	9	2		V31 L1	10	277
HPV31	Ll	10	2	HP'	V31 L1	11	277
HPV31		9	149		V31 L1	10	12
HPV31		11	149		V31 L1	11	12
HPV31	L1	9	299		V31 L1	10	364
HPV31		9	424		V31 L1	9	250
HPV31	L1	10	424		V31 L1	8	445
HPV31	L1	9	283		V31 L1	11	27
HPV31		9	23		V31 L2	9	24
HPV31	L1	11	23		V31 L2	10	24
HPV31		8	340		V31 L2	8	143
HPV31		9	340		V31 L2	10	143
HPV31		10	340		V31 L2	8	281
HPV31	L1	11	340		V31 L2	8	286
HPV31	L1	11	492		V31 L2	9	286
HPV31		11	290		V31 L2	9	367
HPV31		11	344		V31 L2	10	367
HPV31		8	194		V31 L2	11	367
HPV31		9	194		V31 L2	10	15
HPV31		11	194		V31 L2	11	15
HPV31		8	271		V31 L2	8	226
HPV31		10	212		V31 L2	11	226
HPV31		8	284		V31 L2	9	135
HPV31		10	286		V31 L2	10	135
HPV31		11	246		V31 L2	11	135 342
HPV31		8	383		V31 L2	11	376
HPV31		10	383		V31 L2 V31 L2	10	376
HPV31		9	96 96		V31 L2 V31 L2	8	382
HPV31		11 9	494		V31 L2 V31 L2	10	382
HPV31 HPV31		8	494		V31 L2 V31 L2	11	382
					V31 L2	8	133
HPV31 HPV31		11 9	408 .		V31 L2	9	133
HPV31		11	337		V31 L2	11	133
HPV31		10	493		V31 L2	9	278
HPV31		8	267		V31 L2	10	278
HPV31		10	267		V31 L2	11	278
HPV31		9	44		V31 L2	10	400
HPV31		9	333		V31 L2	8	322
HPV31		11	333		V31 L2	9	354
HPV31		9	10		V31 L2	10	354
HPV31		8	239		V31 L2	9	43
HPV31		9	239		V31 L2	11	43
HPV31		8	195		V31 L2	8	358
HPV31		10	195		V31 L2	10	358
HPV31		10	28		V31 L2	11	358
HPV31		11	422		V31 L2	8	364
HPV31		10	332		V31 L2	9	139
HPV31		8	334		V31 L2	10	139
HPV31		10	334		V31 L2	8	116
HPV31		8	62		V31 L2	9	31
HPV31		8	21		V31 L2	10	31
HPV31		9	21	HP	V31 L2	11	31
	-						

157

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				-		
HPV31	L2	9	408	HPV31 L2	10	245
HPV31	L2	11	408	HPV31 L2	11	245
HPV31		10	84	HPV31 L2	8	114
HPV31		8	190	HPV31 L2	10	114
HPV31		9	190	HPV31 L2	10	105
HPV31		11	190	HPV31 L2	11	105
HPV31		9	334	HPV31 L2	9	197
HPV31		10	334	HPV31 L2	10	197
HPV31		11	334	HPV31 L2	10	23
HPV31		8	171 253	HPV31 L2 HPV31 L2	11	23 225
HPV31 HPV31		9 10	196	HPV31 L2	9	225
HPV31		11	196	HPV31 L2	8	35
HPV31		11	276	HPV31 L2	11	35
HPV31		8	237	HPV31 L2	10	242
HPV31		9	237	HPV31 L2	10	302
HPV31		10	237	HPV31 L2	11	302
HPV31		11	158	HPV31 L2	8	231
HPV31		8	459	HPV31 L2	10	231
HPV31	L2	8	361	HPV31 L2	8	423
HPV31	L2	11 .	361	HPV31 L2	10	423
HPV31	L2	9	433	HPV31 L2	8	244
HPV31	L2	11	118	HPV31 L2	11	244
HPV31		10	314	HPV31 L2	8	176
HPV31		9	339	HPV31 L2	10	176
HPV31		10	339	HPV31 L2	9	177
HPV31		8	310	HPV31 L2	9	164
HPV31		8	59	HPV31 L2	8	287 108
HPV31 HPV31		9	113 113	HPV31 L2 HPV31 L2	8 10	108
HPV31		11	57	HPV31 L2	9	447
HPV31		9	351	HPV31 L2	8	335
HPV31		10	221	HPV31 L2	9	335
HPV31		8	26	HPV31 L2	10	335
HPV31		11	26	HPV31 L2	11	335
HPV31	L2	9	65	HPV31 L2	11	256
HPV31	L2	11	65	HPV31 L2	9	269
HPV31	L2	9	52	HPV31 L2	11	269
HPV31	L2	8	213	HPV31 L2	8	204
HPV31		10	213	HPV31 L2	11	204
HPV31		11	413	HPV31 L2	8	390
HPV31		9	175	HPV31 L2	8	292 370
HPV31		11	175	HPV31 L2	8 11	370
HPV31		8	38	HPV31 L2 HPV31 L2	8,	169
HPV31 HPV31		9 11	38 41	HPV31 L2	9	169
HPV31		8	280	HPV31 L2	10	169
HPV31		9	280	HPV31 L2	8	328
HPV31		8	270	HPV31 L2	11	328
HPV31		10	270	HPV31 L2	9	142
HPV31		11	270	HPV31 L2	11	142
HPV31		8	134	HPV31 L2	9	285
HPV31		10	134	HPV31 L2	10	285
HPV31		11	134	HPV31 L2	9	120
HPV31	L2	8	279	HPV31 L2	10	120
HPV31	L2	9	279	HPV31 L2	10	217
HPV31		10	279	HPV31 L2	11	217
HPV31		9	144	HPV31 L2	10	366
HPV31		9	45	HPV31 L2	11	366
HPV31	L2	10	205	HPV31 L2	8	250

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				-5			
HPV31	L2	9	410	HPV31 L	2	10	298
HPV31	L2	8	402	HPV31 L	2	9	69
HPV31	L2	10	402	HPV31 L	2	8	9
HPV31	L2	9	210	HPV31 L	2	10	9
HPV31	L2	11	210	HPV31 L	-2	8	306
HPV31	L2	8	122	HPV31 L	.2	10	306
HPV31	L2	8	88	HPV31 L		8	316
HPV31	L2	11	88	HPV31 L	12	11	316
HPV31	L2	9	422	HPV31 L		9	454
HPV31	L2	11	422	HPV31 L		11	454
HPV31	L2	9	100	HPV31 L		8	239
HPV31	L2	10	100	HPV31 L		8	14
HPV31	L2	8	337	HPV31 L		11	14
HPV31	L2	9	337	HPV31 L		8	341
HPV31	L2	11	337	HPV31 L		9	381
HPV31		8	394	HPV31 L		11	381
HPV31		10	394	HPV31 L		8	384
HPV31		8	74	HPV31 L		9	384
HPV31		9	74	HPV31 L		10	384
HPV31		9	192	HPV31 L		8	94
HPV31		10	235	HPV31 L		9	332
HPV31		11	235	HPV31 L		11	332
HPV31		8	156	HPV31 L		11	431
HPV31		9	156	HPV31 L		9	325
HPV31		8	388	HPV31 L		11	325
HPV31		10	388	HPV31 L		8	86
HPV31		10	167	HPV31 L		10	86
HPV31		11	167	HPV31 L		10	182
HPV31		9	415	HPV31 L		11	104
HPV31		10	415	HPV31 L		8	107
HPV31		8	425	HPV31 L		9	107
HPV31		10	425	HPV31 L		11	107
HPV31		8	127	HPV31 L		11 9	260 50
HPV31		9	127	HPV31 L		11	50
HPV31		10	127	HPV31 L		9	374
HPV31		11	127 97	HPV31 L		10	374
HPV31 HPV31		9 10	97	HPV31 L		8	396
HPV31		10	92	HPV31 L		9	151
HPV31		8	44	HPV31 L		8	184
HPV31		10	44	HPV31 L		10	184
HPV31		9	243	HPV31 L		8	6
HPV31		8	17		.2	10	6
HPV31		9	17	HPV31 L		11	6
HPV31		11	17	HPV31 L		10	346
HPV31		9	228	HPV31 L		8	199
HPV31		11	228	HPV31 L		11	199
HPV31		8	20	HPV31 L		11	208
HPV31		9	303	HPV31 L		10	76
HPV31		10	303	HPV31 L		9	379
HPV31		11	303	HPV31 L		11	379
HPV31		8	417		.2	8	80
HPV31		10	417	HPV31 L	2	10	80
HPV31		11	417	HPV31 L	2	9	162
HPV31		8	229	HPV31 L	2	11	162
HPV31		10	229	HPV31 L	.2	9	149
HPV31		10	12	HPV31 L	2	11	149
HPV31		8	219	HPV31 L	.2	8	137
HPV31		9	219	HPV31 I	12	9	137
HPV31		8	298	HPV31 L	12	11	137

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			•	• •		
HPV31	L2	8	375	HPV33 E1	8	65
HPV31		9	375	HPV33 E1	9	532
HPV31		11	375	HPV33 E1	11	532
HPV31		9	347	HPV33 E1	8	84
HPV31		11	347	HPV33 E1	10	546
HPV31		8	304	HPV33 E1	11	546
HPV31		9	304	HPV33 E1	8	311
HPV31		10	304	HPV33 E1 HPV33 E1	9	311
HPV31 HPV31		9 10	16 16	HPV33 E1	11	318
HPV31		10	227	HPV33 E1	10	373
HPV31		8	416	HPV33 E1	11	373
HPV31		9	416	HPV33 E1	10	81
HPV31		11	416	HPV33 E1	11	81
HPV31		8	136	HPV33 E1	11	22
HPV31		9	136	HPV33 E1	8	83
HPV31	L2	10	136	HPV33 E1	9	83
HPV31	L2	8	39	HPV33 E1	8	310
HPV31		8	140	HPV33 E1	9	310
HPV31	L2	9	140	HPV33 E1	10	310
HPV31	L2	11	140	HPV33 E1	11	230
HPV31		9	426	HPV33 E1	8	259
HPV31		8	128	HPV33 E1	9	259
HPV31		9	128	HPV33 E1	10	259
HPV31		10	128	HPV33 E1	11	259
HPV31		11	128	HPV33 E1 HPV33 E1	8 10	465 465
HPV31		9	344	HPV33 E1	9	297
HPV31 HPV31		10 11	343 391	HPV33 E1	11	226
HPV31		10	362	HPV33 E1	9	14
HPV31		8	254	HPV33 E1	10	14
HPV31		10	392	HPV33 E1	11	14
HPV31	L2	9	81	HPV33 E1	8	118
HPV31	L2	9	232	HPV33 E1	11	118
HPV31	L2	.8	32	HPV33 E1	10	494
HPV31	L2	9	32	HPV33 E1	10	508
HPV31		10	32	HPV33 E1	11	508
HPV31		11	32	HPV33 E1	9	367
HPV31		8	163	HPV33 E1	10	367 46
HPV31		10	163	HPV33 E1 HPV33 E1	10 8	78
HPV31 HPV31		9	377 377	HPV33 E1	9	349
HPV31		11	147	HPV33 E1	8	62
HPV31		8	356	HPV33 E1	11	62
HPV31		10	356	HPV33 E1	8	541
HPV31		8	440	HPV33 E1	10	541
HPV31		9	440	HPV33 E1	10	324
HPV31	L2	10	446	HPV33 E1	8	516
HPV31	L2	9	19	HPV33 E1	10	516
HPV31	L2	10	72	HPV33 E1	9	64
HPV31		11	72	HPV33 E1	8	21
HPV31		8	386	HPV33 E1	8	206
HPV31		10	386	HPV33 E1	10	206
HPV33		11	382	HPV33 E1	11	206
HPV33		8	90	HPV33 E1	10	537 537
HPV33		11	90	HPV33 E1 HPV33 E1	11	186
HPV33		9	96 96	HPV33 E1	10	127
HPV33 HPV33		10 10	383	HPV33 E1	8	361
HPV33		9	104	HPV33 E1	9	361
UL 422	EI	,	104		-	301

Table VIII
HLA-A2 Supermotif-Bearing Peptides

HPV33	E1	11	214	HPV33 E1	11	197
HPV33		11	352	HPV33 E1	8	510
HPV33		8	38	HPV33 E1	9	510
HPV33		10	38	HPV33 E1	11	510
HPV33		11	38	HPV33 E1	8	393
HPV33		11	295	HPV33 E1	9	393
HPV33		10	173	HPV33 E1	9	285
HPV33		11	173	HPV33 E1	8	304
HPV33		9	139	HPV33 E1	9	304
HPV33		10	19	HPV33 E1	8	412
HPV33		8	137	HPV33 E1	9	249
HPV33		11	137	HPV33 E1	11	425 223
HPV33		8	169 89	HPV33 E1 HPV33 E1	9 . 10	223
HPV33				HPV33 E1	9	245
HPV33 HPV33	E1	9 10	89 50	HPV33 E1	11	245
HPV33		9	449	HPV33 E1	8	375
HPV33		10	486	HPV33 E1	9	375
HPV33		11	456	HPV33 E1	10	375
HPV33		8	385	HPV33 E1	8	467
HPV33		10	385	HPV33 E1	9	247
HPV33		11	385	HPV33 E1	11	247
HPV33		10	451	HPV33 E1	9	483
HPV33	E1	8	265	HPV33 E1	10	483
HPV33		10	399	HPV33 E1	11	271
HPV33		8	459	HPV33 E1	9	47
HPV33		9	459	HPV33 E1	9	555
HPV33	E1	10	459	HPV33 E1	11	555
HPV33	E1	8	209	HPV33 E1	10	438
HPV33	E1	10	235	HPV33 E1	11	438
HPV33	E1	10	11	HPV33 E1	9	290
HPV33	E1	9	512	HPV33 E1	10	290
HPV33	E1	8	480	HPV33 E1	11	290
HPV33		11	416	HPV33 E1	8	556
HPV33		8	44	HPV33 E1	10	556
HPV33		9	564	HPV33 E1	8	286
HPV33		10	6	HPV33 E1	11	286
HPV33		10	327	HPV33 E1	10	257 257
HPV33		11	327	HPV33 E1	11	184
HPV33	E1	8	163	HPV33 E1	8 9	339
HPV33		11	256	HPV33 E1	9	503
HPV33 HPV33	E1 E1	9	368 368	HPV33 E1	11	503
HPV33	E1	8	200	HPV33 E1	8	260
HPV33		11	200	HPV33 E1	9	260
HPV33	E1	9	400	HPV33 E1	10	260
HPV33		8	59	HPV33 E1	11	260
HPV33		10	59	HPV33 E1	8	362
HPV33	E1	11	59	HPV33 E1	9	557
HPV33	E1	8	72	HPV33 E1	10	281
HPV33		10	72	HPV33 E1	11	281
HPV33		11	72	HPV33 E1	9	547
HPV33		8	484	HPV33 E1	10	547
HPV33		9	484	HPV33 E1	11	547
HPV33		8	419	HPV33 E1	10	215
HPV33		10	231	HPV33 E1	10	1
HPV33	E1	11	231	HPV33 E1	8	513
HPV33	E1	8	394	HPV33 E1	11	513
HPV33	E1	10	435	HPV33 E1	9	466
HPV33	E1	9	407	HPV33 E1	8	298

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 S	upermotif-Bearin	g Peptide	:s		
HPV33	R1	10	353		HPV33	E1	8	280
HPV33		11	353		HPV33		11	280
HPV33		11	562		HPV33		10	316
HPV33		8	531		HPV33		11	109
HPV33		10	531		HPV33		8	608
HPV33		11	80		HPV33		9	608
HPV33		8	443		HPV33		10	95
HPV33		10	443		HPV33		11	95
HPV33		8	346		HPV33		9	634
HPV33		9	346		HPV33		8	539
HPV33		9	199		HPV33		9	539
HPV33		9	71		HPV33		10	539
HPV33		11	71		HPV33		9	111
HPV33		8	321		HPV33		8	292
HPV33		9	321		HPV33		9	292
HPV33		9	31		HPV33		8	58
HPV33		10	31		HPV33		9	58
HPV33		9	627		HPV33		11	58
HPV33		8	289		HPV33		10	482
HPV33		10	289		HPV33		11	482
HPV33		11	289		HPV33		11	243
HPV33		10	155		HPV33		9	252
HPV33		9	135		HPV33		10	252
HPV33		10	135		HPV33		8	54
HPV33		9	473		HPV33		10	54
HPV33		11	473		HPV33		11	390
HPV33		8	175		HPV33		8	149
HPV33		9	175		HPV33		11	149
HPV33		10	175		HPV33		8	93
HPV33		8	189		HPV33		9	93
HPV33		10	189		HPV33		11	307
HPV33		10	181		HPV33		11	629
HPV33		11	181		HPV33		8	239
HPV33		11	471		HPV33		9	239
HPV33		11	519		HPV33		10	239
HPV33		11	434		HPV33		9	39
HPV33		8	554		HPV33		10	39
HPV33		10	554		HPV33	E1	11	447
HPV33		9	505		HPV33	E1	9	317
HPV33		10	505		HPV33	E1	8	183
HPV33		9	60		HPV33	E1	9	183
HPV33		10	60		HPV33	E1	8	140
HPV33	E1	10	391		HPV33	E1	9	328
HPV33	E1	11	391		HPV33	E1	10	328
HPV33		9	374		HPV33	E1	11	328
HPV33	E1	10	374		HPV33	E1	9	282
HPV33	E1	11	374		HPV33	E1	10	282
HPV33	E1	8	549		HPV33	E1	11	337
HPV33	E1	9	549		HPV33	E1	8	240
HPV33	E1	10	549		HPV33	E1	9	240
HPV33	E1	8	437		HPV33	E1	8	283
HPV33		11	437		HPV33	E1	9	283
HPV33		10	145		HPV33	E1	11	283
HPV33		10	308		HPV33	E1	11	299
HPV33		11	308		HPV33	E1	10	23
HPV33		9	146		HPV33		9	190
HPV33		11	146		HPV33	E1	11	190
HPV33	E1	8	103		HPV33	E1	8	246
HPV33		10	103		HPV33	E1	10	246
HPV33		11	545		HPV33	E1	10	338

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				-				
HPV33	E1	9	325		HPV33	E2	9	136
HPV33	E1	8	548		HPV33	E2	11	136
HPV33	E1	9	548		HPV33	E2	8	74
HPV33	E1	10	548		HPV33	E2	9	74
HPV33	E1	11	548		HPV33	E2	9	298
HPV33	E1	9	436		HPV33	E2	8	328
HPV33	E1	11	144		HPV33	E2	10	328
HPV33	E1	9	354		HPV33	E2	11	328
HPV33	E1	10	354		HPV33	E2	10	80
HPV33		9	182		HPV33	E2	8	185
	E1	10	182		HPV33	E2	9	185
HPV33	E1	9	517		HPV33	E2	10	185
HPV33	E1	11	100		HPV33	E2	10	334
HPV33	E1	8	356		HPV33	E2	11	334
	E1	8	332		HPV33	E2	8	70
	E1		418		HPV33	E2	9	70
	E1		502		HPV33	E2	10	70
	E1	10	502		HPV33	E2	9	325
HPV33	E1	8	522		HPV33	E2	11	325
HPV33	E1	11	522		HPV33	E2	8	319
HPV33	E1	11	372		HPV33	E2	9	319
HPV33	E1	9	524		HPV33	E2	11	156
HPV33	E1	10	524		HPV33	E2	8	190
	E1	8	571		HPV33	E2	10	190
HPV33		11	528		HPV33	E2	8	336
HPV33		8	441		HPV33	E2	9	336
HPV33	E1	10	441		HPV33	E2	11	336
HPV33	E1	8	254		HPV33	E2	10	53
HPV33	E2	9	223		HPV33	E2	11	53
HPV33	E2	11	223		HPV33	E2	9	278
HPV33	E2	8	224		HPV33	E2	8	56
HPV33	E2	10	224		HPV33	E2	9	56
HPV33	E2	11	224		HPV33	E2	8	187
HPV33	E2	9	175		HPV33	E2	11	187
	E2	9	249		HPV33	E2	8	139
	E2		249		HPV33	E2	9	139
	E2	11	249		HPV33	E2	10	139
	E2	9	41		HPV33	E2	11	139
	E2	10	237		HPV33	E2	10	15
	E2	10	258		HPV33	E2	11	276
	E2	11	258		HPV33	E2	8	320
	E2	9	10		HPV33	E2	8	68
	E2	10	245		HPV33	E2	10	68
HPV33		11	245		HPV33	E2	11	68
HPV33		10	40		HPV33	E2	8	14
	E2	8	145		HPV33	E2 E2	11	14 339
HPV33		11	145		HPV33	E2	8 10	339
	E2	8	261		HPV33 HPV33	E2	8	242
	E2	8	174		HPV33	E2 .	9	242
	E2 E2	10	174		HPV33	E2 E2	8	34
			25		HPV33	E2	9	34
	E2 E2	8 10	17		HPV33	E2	10	34
	62 E2	9	17 235		HPV33	E2	8	112
			143		HPV33	E2	8	47
	E2 E2	10 9	143 232		HPV33	E2 E2	10	264
					HPV33	E2	11	264
HPV33 I	E2	10 11	232 20		HPV33	E2	8	23
	E2 E2	8				E2	8	66
		9	3		HPV33		10	66
HPV33	52	,	3		11E 422		10	-

163

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TEST TIE Supermout De	arms repude	•		
HPV33	E2	9	180	HPV33	E2	9	191
HPV33	E2	10	151	HPV33	E2	8	57
HPV33	E2	11	165	HPV33	E2	11	57
HPV33		10	63	HPV33	E2	8	292
HPV33		11	63	HPV33	E2	9	7
HPV33		8	35	HPV33	E2	10	7
HPV33		9 .	35	HPV33	E2	9	37
HPV33		11	35	HPV33	E2	10	37
HPV33		8	62	HPV33		10	253
HPV33		11	62	HPV33	E2	8	266
HPV33		8	42	HPV33	E2	9	266
HPV33		8	75	HPV33		11	266
HPV33		8	94	HPV33	E2	11	5
HPV33		10	94	HPV33		9	198
HPV33		10	240	HPV33		10	198
HPV33		11	240	HPV33	E2	9	285
HPV33		9	147	HPV33		9	61
HPV33		11	147	HPV33		9	302
HPV33		8	9	HPV33	E2	8	28
HPV33		10	9	HPV33		9	28
HPV33		8	202	HPV33		11	28
HPV33		9	202	HPV33	E2	8	90
HPV33		11	202	HPV33	E2	10	90
HPV33		11	127	HPV33	E2	9	85
HPV33		8	272	HPV33	E2	10	85
HPV33		8	230	HPV33	E2	10	88
HPV33		11	230	HPV33		8	205
HPV33		8	248	HPV33	E2	10	205
HPV33	E2	10	248	HPV33	E2	10	45
HPV33		11	248	HPV33	E2	8	236
HPV33		8	239	HPV33	E2	11	236
HPV33		11	239	HPV33	E2	9	254
HPV33	E2	11	221	HPV33	E2	11	257
HPV33	E2	9	196	HPV33	E2	9	93
HPV33	E2	11	196	HPV33	E2	11	93
HPV33	E2	10	342	HPV33	E2	9	81
HPV33	E2	11	342	HPV33	E2	10	128
HPV33	E2	10	222	HPV33		10	146
HPV33	E2	8	29	HPV33	E2	8	181
HPV33	E2	10	29	HPV33	E2	8	233
HPV33	E2	8	345	HPV33		9	233
HPV33	E2	9	345	HPV33		11	233
HPV33	E2	8	203	HPV33		9	206
HPV33	E2	10	203	HPV33		8	267
HPV33		9	332	HPV33		10	267
HPV33		11	48	HPV33		11	267
HPV33		11	182	HPV33		8	337
HPV33		8	331	HPV33		10	337
HPV33		10	331	HPV33		9	343
HPV33		8	330	HPV33		10	343
HPV33		9	330	HPV33		11	343
HPV33		11	330	HPV33		11	118
HPV33		9	329	HPV33		8	72
HPV33		10	329	HPV33		10	72
HPV33		9	95	HPV33		11	72
HPV33		11	213	HPV33		8	192
HPV33		8	96	HPV33		8	11
HPV33		8	71	HPV33		11	11
HPV33		9	71	HPV33		8	344
HPV33	E2	11	71	HPV33	E2	9	344

Table VIII HLA-A2 Supermotif-Bearing Peptides

			nLA-A2 Supermont-Beam	ng repud	C3		
HPV33	E2	10	344	HPV33	E5	11	1
HPV33		10	119	HPV33	E5	8	61
HPV33	E2	11	119	HPV33	E5	11	61
HPV33	E2	8	326	HPV33	E5	8	21
HPV33	E2	10	326	HPV33	E5	10	21
HPV33	E2	11	323	HPV33	E5	9	46
HPV33	E2	8	148	HPV33	E5 ·	10	46
HPV33	E2	10	148	HPV33	E5	9	60
HPV33		10	58	HPV33	E5	8	25
HPV33	E2	8	92	HPV33	E5	9	25
HPV33		10	92	HPV33	E5	10	25
HPV33	E2	8	159	HPV33	E5	11	25
HPV33	E2	9	138	HPV33	E5	8	16
HPV33		10	138	HPV33	E5	9	16
HPV33	E2	11	138	HPV33	E5	11	16
HPV33	E2	11	44	HPV33	E5	8	27
HPV33	E2	9	131	HPV33	E5	9	27
HPV33	E2	10	131	HPV33	E5	10	27
HPV33	E5	9	63	HPV33	E5	11	27
HPV33	E5	10	63	HPV33	E5	8	6
HPV33	E5	9	14	HPV33	E5	10	6
HPV33	E5	10	14	HPV33	Ė5	8	36
HPV33	E5	11	14	HPV33	E5	8	34
HPV33	E5	9	9	HPV33	E5	10	34
HPV33	E5	10	9	HPV33	E5	8	31
HPV33	E5	11	9	HPV33	E5	9	31
HPV33	E5	8	12	HPV33	E5	11	31
HPV33	E5	11	12	HPV33	E5	8	40
HPV33	E5	9	56	HPV33	E5 -	10	40
HPV33	E5	8	3	HPV33		8	29
HPV33	E5	9	3	HPV33	E5	9	29
HPV33	E5	11	3	HPV33	E5	10	29
HPV33		8	42	HPV33		11	29
HPV33	E5	9	5	HPV33	E5	9	53
HPV33	E5	11	5	HPV33		10	53
HPV33	E5 -	8	10	HPV33	E5	11	58
HPV33	E5	9	10	HPV33	E6	11	137
HPV33	E5	10	10	HPV33	E6	9	18
HPV33		8	23	HPV33	E6	11	18
HPV33		10	23	HPV33	E6	8	103
HPV33		11	23	HPV33	E6	8	66
HPV33		8	48	HPV33	E6	8	16
HPV33		10	48	HPV33	E6	11	16
HPV33		9	22	HPV33	E6	8	30
HPV33		11	22	HPV33	E6	8	14
HPV33		8	54	HPV33	E6	9	14
HPV33		9	54	HPV33	E6	10	14
HPV33		11	54	HPV33	E6	9	120
HPV33		8	17	HPV33	E6	8	4
	E5	10	17	HPV33	E6	9	4 98
HPV33		11	37	HPV33	E6	8	27
HPV33	E5	9	18	HPV33	E6	11	89
HPV33		11	18	HPV33	E6	8	89
	E5	8	32	HPV33	E6	11	
HPV33	E5	10	32	HPV33	E6	9	20
HPV33		10	38	HPV33	E6	8	41
HPV33		9	35	HPV33	E6	10	41
HPV33		9	33	HPV33		11	41
HPV33		11	33	HPV33	E6	8	45
HPV33	E5	10	1	HPV33	E6	9	45

Table VIII HLA-A2 Supermotif-Bearing Peptides

			•			
HPV33	E6	10	2	HPV33 E7	8	88
HPV33		11	2	HPV33 E7	8	62
HPV33		10	61	HPV33 E7		62
HPV33		11	118	HPV33 E7	8	47
HPV33		10	64	HPV33 E7	9	47
HPV33		11	100	HPV33 E7	10	47
HPV33		10	28	HPV33 E7	10	19
HPV33		10	37	HPV33 E7	8	6
HPV33		11	127	HPV33 E7	10	6
	E6	11	86	HPV33 E7 HPV33 E7	11 9	44 81
HPV33 HPV33		11	109 95	HPV33 E7	10	81
HPV33		11	95	HPV33 E7	8	80
HPV33		11	36	HPV33 E7	10	80
	E6	8	112	HPV33 E7	11	80
HPV33		10	112	HPV33 E7	8	66
HPV33		10	17	HPV33 E7	11	66
HPV33	E6	10	90 -	HPV33 E7	8	77.
HPV33	E6	11	90	HPV33 E7	10	77
HPV33	E6	9	10	HPV33 E7	11	77
HPV33		10	10	HPV33 E7	8	71
HPV33	E6	10	82	HPV33 E7	9	71
HPV33		11	82	HPV33 E7	8	49
HPV33	E6	10	22	HPV33 E7	8	72
HPV33		10	87	HPV33 E7	11	72
HPV33		8	11	HPV33 E7	9	78
HPV33		9	11	HPV33 E7	10 9	78 ·
HPV33	E6	11	11	HPV33 E7 HPV33 E7	10	63
HPV33 HPV33	E6	8	21 .	HPV33 E7	11	63
HPV33		9	91	HPV33 E7	8	86
HPV33	E6	10	91	HPV33 E7	10	86
HPV33	E6	11	91	HPV33 E7	9	64
	E6	11	52	HPV33 E7	10	64
HPV33	E7	10	45	HPV33 E7	9	12
HPV33		11	45	HPV33 E7	11	12
HPV33	E7	8	48	HPV33 E7	9	55
HPV33	E7	9	48	HPV33 E7	10	55
HPV33		9	68	HPV33 E7	11	55
HPV33		11	68	HPV33 E7	8	53
HPV33		8	75	HPV33 E7	11	53
	E7	9	75	HPV33 E7	10	11
HPV33	E7	10	75	HPV33 L1 HPV33 L1	10	179 482
HPV33	E7	8	21	HPV33 L1	8 11	482
HPV33 HPV33	E7 E7	9	14 37	HPV33 L1	8	424
HPV33	E7	8	43	HPV33 L1	8	316
HPV33	E7	9	85	HPV33 L1	8	9
HPV33		11	85	HPV33 L1	10	9
HPV33	E7	9	59	HPV33 L1	9	44
	E7	11	59	HPV33 L1	8	270
HPV33	E7	8	79	HPV33 L1	9	158
HPV33	E7	9	79	HPV33 L1	9	147
HPV33	E7	11	79	HPV33 L1	11	147
HPV33	E7	10	54	HPV33 L1	9	207
HPV33	E7	11	54	HPV33 L1	9	345
HPV33	E7	8	82	HPV33 L1	10	396
HPV33	E7	9	82	HPV33 L1	11	396
HPV33	E7	8	83	HPV33 L1	8	449
HPV33	E7	11	83	HPV33 L1	10	449

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bearin	ng Peptides		
HPV33	T.1	8	370	HPV33 L1	10	115
HPV33		11	370	HPV33 L1	8	391
		9	274	HPV33 L1	10	365
HPV33		10	274	HPV33 L1	8	194
HPV33		11	274	HPV33 L1	10	194
HPV33		10	199	HPV33 L1	9	397
HPV33		10	459	HPV33 L1	10	397
HPV33		11	459	HPV33 L1	9	286
HPV33		11	202	HPV33 L1	9	474
HPV33		8	95	HPV33 L1	10	474
HPV33		10	95	HPV33 L1	8	478
HPV33		9	88	HPV33 L1	10	478
HPV33		8	335	HPV33 L1	10	60
HPV33		9	335	HPV33 L1	11	236
HPV33		10	335	HPV33 L1	8	153
HPV33		10	415	HPV33 L1	10	211
HPV33		11	415	HPV33 L1	9	65
HPV33		10	219	HPV33 L1	8	379
HPV33		9	8	HPV33 L1	9	379
HPV33		11	8	HPV33 L1	10	379
HPV33		9	269	HPV33 L1	9	20
HPV33		8	107	HPV33 L1	10	43
HPV33		10	107	HPV33 L1	11	190
HPV33		10	447	HPV33 L1	11	42
HPV33		8	385	HPV33 L1	8	159
HPV33		9	385	HPV33 L1	8	123
HPV33		9	467	HPV33 L1	10	123
HPV33		9	249	HPV33 L1	8	468
HPV33	L1	8	375-	HPV33 L1	9	61
HPV33	L1	9	375	HPV33 L1	11	469
HPV33	L1	10	375	HPV33 L1	8	213
HPV33	L1	8	373	HPV33 L1	10	213
HPV33	L1	9	373	HPV33 L1	8	413
HPV33	L1	10	373	HPV33 L1	9	413
HPV33	L1	11	373	HPV33 L1	10	371
HPV33	L1	9 .	256	HPV33 L1	11	371
HPV33	L1	11	256	HPV33 L1	11	313
HPV33	L1	8	419	HPV33 L1	8	69
HPV33		8	330	HPV33 L1	10	69
HPV33		11	330	HPV33 L1	9	382
HPV33		8	141	HPV33 L1	11	382
HPV33		10	141	HPV33 L1	9	405
HPV33		8	322	HPV33 L1	8	62
HPV33		10	322	HPV33 L1	8 10	99 99
HPV33		11	322			342
HPV33		8	117	HPV33 L1	8	237
HPV33		10	105	HPV33 L1 HPV33 L1	10	387
HPV33		8	472	HPV33 L1	11 9	200
HPV33		11	472	HPV33 L1	8	299
HPV33		9	68	HPV33 L1	11	178
HPV33		11	68	HPV33 L1	10	57
HPV33		8 10	404	HPV33 L1	8	341
HPV33				HPV33 L1	9	341
HPV33		11	138	HPV33 L1	8	327
HPV33		8	111	HPV33 L1	10	327
HPV33		11	265	HPV33 L1	11	327
HPV33		10	281 173	HPV33 L1	9	192
HPV33		8		HPV33 L1	10	192
HPV33		10	173	HPV33 L1	8	181
HPV33	ΠŢ	9	115	III 433 DT	o	101

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			•		•		
HPV33	L1	10	181	HPV	733 L1	9	23
HPV33	L1	11	181	HPV	/33 L1	11	23
HPV33	L1	8	221	HPV	/33 L1	8	339
HPV33	L1	8	187	HPV	/33 L1	10	339
HPV33	L1	9	187	HPV	733 L1	11	339
HPV33	L1	10	439	HPV	733 L1	8	2
HPV33	L1	8	462	HPV	/33 L1	9	2
HPV33	L1	10	462			10	2
HPV33	L1	11	113				383
HPV33	L1	9	122				383
HPV33		11	122				383
HPV33		10	165				283
HPV33		11	165				193
HPV33		8	55				193
HPV33		9	55				193
HPV33		8	293				343
HPV33		9	293				266
HPV33		9	484				212
HPV33		8	15				212
HPV33		10	15				381
HPV33		8	17				381
HPV33		10	470				96
HPV33		8	423				96 346
HPV33		9	423				282
HPV33		9	214 214				336
HPV33 HPV33		11	376				336
		9	376				336
HPV33		11	376				430
HPV33		11	156				332
HPV33		9	305				332
HPV33		11	254				10
HPV33		9	328				174
HPV33		10	328			8	386
HPV33		9	481	HPV	/33 L1	8	380
HPV33		8	263	HPV	/33 L1	9	380
HPV33	L1	9	263	HPV	/33 L1	11	380
HPV33	L1	9	315	HPV	/33 L1	11	420
HPV33	L1	9	98	HPV	/33 L1	10	331
HPV33	L1	11	98	HPV	/33 L1	8	333
HPV33	L1	8	30			10	333
HPV33	L1	10	488			11	333
HPV33		9	455			8	21
HPV33		11	289			11	21
HPV33		11	410			8	101
HPV33		8	51			11	401
HPV33		11	51			8	36
HPV33		10	285		/33 L1	9	36
HPV33		11	32			10	36
HPV33		11	245			9	389
HPV33		9	412			10	389
HPV33		10	412		/33 L1	8	276 276
HPV33		9	149		/33 L1	9	276
HPV33		11	149			11	
HPV33		9	298		/33 L1	10	362 12
HPV33		9	422			10	12
HPV33		10	422		/33 L1 /33 L1	11 8	443
HPV33		10	304		/33 L1	8 11	27
HPV33		9	227		/33 L2	8	81.
HPV33	LI	11	227	HP	/33 112	0	01.

WO 01/41799

Table VIII

PCT/US00/33549

			HLA-A2 Supermotif-Bea	ring Peptides		
HPV33	L2	8	140	HPV33 L2	11	201
HPV33	L2	11	82	HPV33 L2	9	117
HPV33	L2	8 .	291	HPV33 L2	10	319
HPV33	L2	9	291	HPV33 L2	10	361
HPV33	L2	8	286	HPV33 L2	10	226
HPV33	L2	9	23	HPV33 L2	8	305
HPV33	L2	10	23	HPV33 L2	8	25
HPV33	L2	11	308	HPV33 L2	11	25
HPV33	L2	10	14	HPV33 L2	8	75
HPV33	L2	11	14	HPV33 L2	9	75
HPV33	L2	8	385	HPV33 L2	11	75
HPV33	L2	10	385	HPV33 L2	10	60
HPV33	L2	9	283	HPV33 L2	9	51
HPV33		10	283	HPV33 L2	11	51
HPV33		11	283	HPV33 L2	11	158
HPV33		9	409	HPV33 L2	10	374
HPV33		11	272	HPV33 L2	10	336
HPV33		8	327	HPV33 L2	8	297
HPV33		11	327	HPV33 L2	11	40
HPV33		9	42	HPV33 L2	8	285
HPV33		11	42	HPV33 L2	9	285 284
HPV33		8	431	HPV33 L2	9	284
HPV33		11	431	HPV33 L2 HPV33 L2	10	284
HPV33		10	264	HPV33 L2	9	44
HPV33 HPV33		10 9	401 350	HPV33 L2	8	152
HPV33		9	136	HPV33 L2	11	152
HPV33		10	95	HPV33 L2	8	292
HPV33		11	95	HPV33 L2	8	331
HPV33		9	369	HPV33 L2	8	104
HPV33		10	30	HPV33 L2	11	104
HPV33		11	30	HPV33 L2	9	433
HPV33		8	130	HPV33 L2	10	433
HPV33		9	130	HPV33 L2	10	22
HPV33		11	130	HPV33 L2	11	22
HPV33	L2	10	364	HPV33 L2	9	248
HPV33	L2	9	115	HPV33 L2	8	311
HPV33	L2	11	115	HPV33 L2	10	311
HPV33		8	344	HPV33 L2	8	34
HPV33	L2	10	344	HPV33 L2	11	34
HPV33		8	341	HPV33 L2	8	236
HPV33		11	341	HPV33 L2	9	236 107
HPV33		9	110	HPV33 L2	8	107
HPV33		11	110	HPV33 L2	10 8	249
HPV33		9	384	HPV33 L2 HPV33 L2	8	266
HPV33		11	384	HPV33 L2	11	266
HPV33		8 11	113	HPV33 L2	8	85
HPV33 HPV33		8	113 181	HPV33 L2	9	85
HPV33		10	181	HPV33 L2	10	85
HPV33		11	281	HPV33 L2	9	345
HPV33		8	242	HPV33 L2	8	243
HPV33		9	242	HPV33 L2	9	243
HPV33		10	242	HPV33 L2	9	377
HPV33		9	268	HPV33 L2	10	377
HPV33		8	460	HPV33 L2	11	377
HPV33		11	163	HPV33 L2	8	195
HPV33		9	440	HPV33 L2	9	195
HPV33		10	440	HPV33 L2	11	195
HPV33		10	201	HPV33 L2	8	397
1	~-	0				

Table VIII HLA-A2 Supermotif-Bearing Peptides

			Tiert He Supermon Dour	ng r opilato		
HPV33	L2	9	397	HPV33 L2	11	233
HPV33	L2	8	231	HPV33 L2	8	19
HPV33	L2	11	231	HPV33 L2	10	153
HPV33	L2	8	391	HPV33 L2	8	234
HPV33	L2	10	143	HPV33 L2	10	234
HPV33	L2	8	209	HPV33 L2	11	234
HPV33	L2	9	174	HPV33 L2	10	11
HPV33	L2	11	255	HPV33 L2	8	321
HPV33	L2	10	240	HPV33 L2	11	321
HPV33	L2	11	240	HPV33 L2	9	224
HPV33	L2	9	139	HPV33 L2	9	68
HPV33	L2	9	290	HPV33 L2	9	388
HPV33	L2	10	290	HPV33 L2	11	388
HPV33	L2	11	172	HPV33 L2	8	303
HPV33		8	275	HPV33 L2	10	303
HPV33	L2	10	275	HPV33 L2	8	134
HPV33		11	275	HPV33 L2	9	134
HPV33		8	73 .	HPV33 L2	11	134
HPV33		9	73	HPV33 L2	8	13
HPV33		10	73	HPV33 L2	11	13
HPV33		11	73	HPV33 L2	9	357
HPV33		8	215	HPV33 L2	11	357
HPV33		9	215	HPV33 L2	10	393
HPV33		11	215	HPV33 L2	10	122
HPV33		8	87	HPV33 L2	11	122
HPV33		10	87	HPV33 L2	9	151
HPV33		11	87	HPV33 L2	9	103
HPV33		10	423	HPV33 L2	9	49
HPV33		11	423	HPV33 L2	11	49
HPV33		8	330	HPV33 L2	9	106
HPV33		9	330	HPV33 L2	11	106
HPV33		9	99	HPV33 L2 HPV33 L2	8	204
HPV33		10	99	HPV33 L2	11	382
HPV33 HPV33		10	413 395	HPV33 L2	11	382
HPV33		10	395	HPV33 L2	9	156
HPV33		11	395	HPV33 L2	8	38
HPV33		9	84	HPV33 L2	10	213
HPV33		10	84	HPV33 L2	11	213
HPV33		11	84	HPV33 L2	8	189
HPV33		9	197	HPV33 L2	10	189
HPV33		8	376	HPV33 L2	9	6
HPV33		10	376	HPV33 L2	10	6
HPV33		11	376	HPV33 L2	11	352
HPV33		10	79	HPV33 L2	9	146
HPV33		8	161	HPV33 L2	10	146
HPV33		9	161	HPV33 L2	9	167
HPV33		8	124	HPV33 L2	11	167
HPV33		9	124	HPV33 L2	9	80
HPV33	L2	10	124	HPV33 L2	9	154
HPV33		9	416	HPV33 L2	11	154
HPV33	L2	11	186	HPV33 L2	10	432
HPV33		8	403	HPV33 L2	11	432
HPV33		10	91	HPV33 L2	10	309
HPV33		8	43	HPV33 L2	9	265
HPV33	L2	10	43	HPV33 L2	9	15
HPV33	L2	8	16	HPV33 L2	10	15
HPV33	L2	9	16	HPV33 L2	10	232
HPV33	L2	11	16	HPV33 L2	9	190
HPV33	L2	9	233	HPV33 L2	11	190

Table VIII
HLA-A2 Supermotif-Bearing Peptides

HPV33	L2	8	137	HPV45	E1	11	373
HPV33		11	137	HPV45		8	40
HPV33		9	386	HPV45		9	40
HPV33		11	386	HPV45	El	10	251
HPV33		9	132	HPV45	E1	9	202
HPV33	L2	10	132	HPV45	E1 .	10	399
HPV33		11	132	HPV45		11	398
HPV33	L2	8	93	HPV45 HPV45		10	465 465
HPV33	L2	9 10	96 96	HPV45	E1	8	259
HPV33 HPV33		9	337	HPV45	E1	9	259
HPV33		11	298	HPV45		10	259
HPV33		10	187	HPV45	El	11	259
HPV33		11	222	HPV45	E1	9	297
	L2	9	31	HPV45	E1	10	390
HPV33		10	31	HPV45	E1	11	390
HPV33	L2	11	31	HPV45	E1	8	226
HPV33	L2	8	168	HPV45	E1	11	226
	L2	10	168	HPV45	E1	10	634
HPV33	L2	8	441	HPV45	E1	11	621
HPV33		9	441	HPV45	E1	8	78
HPV33		11	404	HPV45		8	516
HPV33		8	131		E1 E1	8	206 206
	L2	10	131	HPV45	E1	10 11	206
HPV33 HPV33		11 9	131 92	HPV45	E1	9	614
HPV33		8	434	HPV45	E1	11	614
HPV33		9	434	HPV45	E1	9	349
	L2	8	237	HPV45	E1	9	108
HPV33		9	202	HPV45	E1	11	108
HPV33		10	202	HPV45	E1	8	361
HPV33	L2	8	366	HPV45	E1	9	361
HPV33	L2	10	366	HPV45	E1	11	214
HPV33	L2	8	325	HPV45	E1	9	367
HPV33		10	325	HPV45		10	367
HPV33	L2	9	18	HPV45	E1 E1	8 10	46 46
HPV33		10	71 71	HPV45		11	352
HPV33 HPV45	L2 E1	11	382	HPV45		8	106
HPV45		8	144	HPV45		11	106
HPV45		10	144	HPV45		9	623
HPV45		10	383	HPV45	E1	10	42
HPV45		8	310	HPV45	E1	10	508
HPV45	E1	10	310	HPV45	E1	11	508
HPV45	E1	10	198	HPV45		9	328
HPV45	E1	8	232	HPV45		10	328
HPV45	E1	9	232	HPV45		11	328
HPV45	E1	10	232	HPV45		10	52 30
HPV45	E1	9	532	HPV45		11	30
HPV45 HPV45	E1 E1	9	68 452	HPV45	E1	8	143
HPV45		9	311	HPV45		9	143
HPV45	E1	9	199	HPV45		11	143
HPV45		9	512	HPV45		11	115
HPV45		10	66	HPV45	E1	8	186
HPV45		8	72	HPV45	E1	10	189
HPV45	E1	10	72	HPV45		11	189
HPV45	E1	11	72	HPV45		8	59
HPV45	El	8	408	HPV45		10	59
HPV45	E1	10	373	HPV45	E1	11	59

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TIETT-TE Supermon-Beath	ng r cpudes		
HPV45	E1	8	62	HPV45 E1	11	257
HPV45	E1	11	62	HPV45 E1	11	333
HPV45	E1	9	101	HPV45 E1	8	184
HPV45		9	64	HPV45 E1	10	184
HPV45		10	38	HPV45 E1	9	23
HPV45		11	38	HPV45 E1	10	23
HPV45		11	295	HPV45 E1	10	435
HPV45		8	21	HPV45 E1	11	197
HPV45	E1	11	21	HPV45 E1	8	124
HPV45		8	146	HPV45 E1	9	124
HPV45	E1	10	146	HPV45 E1	11	425
HPV45		9	141	HPV45 E1 HPV45 E1	9	304
HPV45		10 11	141 141	HPV45 E1 HPV45 E1	8	304 245
HPV45		8	74	HPV45 E1	9	245
HPV45		9	74	HPV45 E1	11	245
HPV45		11	74	HPV45 E1	8	223
	E1	10	324	HPV45 E1	9	223
	E1	11	89	HPV45 E1	10	223
HPV45		8	50	HPV45 E1	11	223
HPV45	E1	8	483	HPV45 E1	8	510
HPV45	E1	9 .	483	HPV45 E1	9	510
HPV45	E1	10	483	HPV45 E1	11	510
HPV45	E1	8	446	HPV45 E1	8	375
HPV45	E1	11	456	HPV45 E1	9	375
	E1	8	385	HPV45 E1	10	375
HPV45		11	385	HPV45 E1	8	506
HPV45		10	486	HPV45 E1	9	506
HPV45		9	449	HPV45 E1	10	201
	E1	10	438	HPV45 E1	11	299
	E1	10	19	HPV45 E1 HPV45 E1	9 11	247 247
HPV45	E1	10 9	494 342	HPV45 E1	9 .	290
HPV45		10	626	HPV45 E1	10	290
	E1	11	626	HPV45 E1	11	290
	E1	8	318	HPV45 E1	9	190
HPV45	E1	8	209	HPV45 E1	10	190
	E1	8	286	HPV45 E1	11	190
HPV45	E1	11	286	HPV45 E1	9	547
HPV45	E1	8	480	HPV45 E1	10	547
HPV45	E1	11	480	HPV45 E1	11	547
HPV45	E1	11	630	HPV45 E1	11	271
HPV45		10	11	HPV45 E1	9	556
	E1	8	459	HPV45 E1	8	467
	E1	9	459	HPV45 E1	8	191
HPV45	E1	10	459	HPV45 E1	9	191
		8	443	HPV45 E1	10	191
HPV45 HPV45		10	443	HPV45 E1 HPV45 E1	11	191 487
		11	443 265	HPV45 E1	8	548
		9	71	HPV45 E1	9	548
		11	71		10	548
		11	256		11	548
		8	83	HPV45 E1	8	362
		11		HPV45 E1	8	336
		8	292	HPV45 E1	8	557
		9	292	HPV45 E1	9	281
		10	555	HPV45 E1	10	281
		9	466	HPV45 E1	11	281
		10	257	HPV45 E1	10	215

172

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TIEA-AZ Supermour-Dearn	g r cpud			
HPV45	E1	11	215	HPV45	E1	8	54
HPV45	E1	8	368	HPV45	E1	8	102
HPV45	E1	9	368	HPV45	E1	8	412
HPV45	El	9	231	HPV45	E1	11	80
HPV45	E1	10	231	HPV45	E1	8	148
HPV45	E1	11	231	HPV45	E1	10	451
	E1	8	200	HPV45	E1	9	407
HPV45		11	200	HPV45	E1	11	612
	E1	8	513	HPV45	E1	9	335
HPV45		11	513	HPV45	E1		280
HPV45	E1	8	298	HPV45	E1		280 280
		9	47	HPV45	E1 E1		411
HPV45	E1 E1	.11	47 353	HPV45	E1		316
		10 11	353	HPV45	E1	8	608
HPV45 HPV45	E1	8	154	HPV45	E1	10	575
	E1	11	154	HPV45	E1	10	56
HPV45	E1	11	174	HPV45	E1		56
HPV45	E1	8	531	HPV45	E1	9	539
HPV45	E1	10	531	HPV45	E1	10	539
HPV45	E1	9	321	HPV45	E1	8	183
HPV45	E1	9	473	HPV45	E1	9	183
HPV45	E1	10	152	HPV45	E1	11	183
HPV45	E1	8	177	HPV45	E1	9	117
	E1	9	177	HPV45	E1	11	416
HPV45	E1	11	177		E1		288
HPV45	E1	10	563	HPV45	E1		288
HPV45	E1	11	471	HPV45	E1 E1	10 10	308 104
HPV45 HPV45	E1 E1	8	250 250	HPV45	E1	8	65
HPV45	E1	11	554	HPV45		11	65
HPV45	E1	11	554	HPV45	E1	9	39
HPV45	E1	11	537	HPV45	E1	10	39
	E1	11	434	HPV45	E1	10	22
HPV45		8	505	HPV45	E1	11	22
HPV45	E1	9	505	HPV45	E1	8	246
HPV45	El	10	505	HPV45	E1	10	246
	E1	8	98	HPV45	E1	8	289
	E1	10	98	HPV45	E1	10	289
	E1	10	546	HPV45	E1	11	289
HPV45	E1	11	546	HPV45	E1 E1	9	252 53
HPV45	E1	9	238	HPV45	E1	9	147
HPV45	E1	10 11	238	HPV45	E1	8	224
	E1	9	60	HPV45	E1	9	224
HPV45	E1	10	60	HPV45	E1	10	224
HPV45	El	9	391	HPV45	E1	8	239
HPV45		10	391	HPV45	E1	9	239
HPV45	E1	11	391	HPV45	E1	10	239
HPV45	E1	9	67	HPV45	E1	8	282
HPV45	El	8	192	HPV45	E1	9	282
HPV45	E1	9	192	HPV45	E1	10	282
	E1	10	192	HPV45	E1	8	577
	E1	11	192	HPV45	E1	9	309
HPV45	E1	9	374	HPV45	E1	11	309
HPV45	E1	10	374	HPV45	E1	8	240
HPV45	E1	11	374	HPV45	E1 E1	9	240
HPV45	E1	8	549	HPV45	E1	9	283
HPV45	E1	9	549	HPV45	E1	11	283
HPV45	E1	10	549	TLA42	ii L	11	203

Table VIII
HLA-A2 Supermotif-Bearing Peptides

				ALDIT TIL Daponi	doin Dearing	. op aa			
HPV45	E1	8	511		н	PV45	E2	10	343
HPV45	E1	10	511		н	PV45	E2	11	343
HPV45	E1	9	31		н	PV45	E2	8	192
HPV45	E1	10	31		н	PV45	E2	9	192
HPV45		11	31		н	PV45	E2	11	349
HPV45	E1	10	81		н	PV45	E2	8	50
HPV45	E1	10	230			PV45	E2	9	50
HPV45	E1	11	230			PV45	E2	11	50
HPV45	E1	9	400			PV45	E2	11	334
HPV45	E1	9	436			PV45	E2	9	56
HPV45	E1	8	75			PV45	E2	10	56
HPV45	E1	10	75			PV45	E2	8	150
HPV45	E1	11	75			PV45	E2	10	150
HPV45	E1	9	354			PV45	E2	10	255
	E1	10	354			PV45	E2	9	237
HPV45	E1	9	635			PV45	E2	11	237
	E1	9	576			PV45	E2	10	163
HPV45	E1	8	356			PV45	E2	11	163
HPV45	E1	9	418			PV45	E2	9	225
	E1	8	332			PV45	E2	10	225
HPV45	E1	8	502			PV45	E2	9	295
HPV45	E1	10	502			PV45	E2	9	62
HPV45	E1	11	502			PV45	E2	11	267
HPV45	E1	8	522			PV45	E2	11	293
	E1	11	522			PV45	E2	10	48
HPV45	E1	8	229			PV45	E2	11	48
	E1	11	229			PV45	E2	10	335
HPV45	E1	11	372			PV45	E2	10	146
HPV45		8	571			PV45	E2	11	146
	E1	10	571			PV45	E2	8	57
	E1	8	394			PV45	E2	9	57
HPV45	E1	11	528			PV45	E2	8	219
HPV45	E1	10	441			PV45	E2	10	219
HPV45		9	156			PV45	E2	8	74
HPV45	E2	8	78			PV45	E2	10	74
HPV45	E2	11	78			PV45	E2	11	74
	E2	11	47			PV45	E2	8	77
HPV45		11	89			PV45	E2	9	77
HPV45		10	247			PV45	E2	10	59
	E2	11	247			PV45	E2	9	2
	E2	8	216			PV45	E2	11	154
	E2	11	216			PV45	E2	8	284
	E2	10	305			PV45	E2	10	284
	E2	10	134			PV45	E2	11	284
HPV45		10	158			PV45	E2	В	41
HPV45		11	31			PV45	E2	9	41
	E2	9	102			PV45	E2	11	41
	E2	8	212			PV45	E2	8	100
	E2	9	212			PV45	E2	11	100
HPV45		9	351			PV45	E2	8	223
HPV45		11	351			PV45	E2	11	223
HPV45	E2	9	319			PV45	E2	8	81
HPV45		9	313			PV45	E2	9	81
HPV45	E2	10	80			PV45	E2	9	256
HPV45	E2	9	258			PV45	E2	11	256
HPV45	E2	11	258			PV45	E2	9	336
HPV45	E2 E2	8	148			PV45	E2	8	3
HPV45	E2	9	148			PV45	E2	10	3 69
HPV45	E2	10	148			PV45	E2	11	69
	E2	8				PV45	E2	8	347
mrv45	E-2	0	343		п	E 4-8 3	62		34/

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TIETE-TE Supermour Dear	ng repudes		
HPV45	E2	9	347	HPV45 E2	8	354
HPV45		8	332	HPV45 E2	10	354
HPV45		8	265	HPV45 E2	9	99
HPV45		9	265	HPV45 E2	10	217
HPV45		8	289	HPV45 E2	8	213
HPV45		11	189	HPV45 E2	11	213
HPV45		9	198	HPV45 E2	8	337
HPV45		11	246	HPV45 E2	8	199
HPV45		9	67	HPV45 E2	11	199
HPV45		8	360	HPV45 E2	8	359
HPV45		9	360	HPV45 E2	9	359
HPV45		8	35	HPV45 E2	10	359
HPV45		10	35	HPV45 E2	9	344
HPV45		9	218	HPV45 E2	10	344
HPV45		11	218	HPV45 E2	11	344
HPV45		8	40	HPV45 E2	8	193
HPV45		9	40	HPV45 E2	9	353
HPV45		10	40	HPV45 E2	11	353
			222	HPV45 E2	11	338
HPV45		9	82	HPV45 E2	8	352
HPV45		11	4	HPV45 E2	10	352
			63	HPV45 E2	9	138
HPV45		9	43	HPV45 E2	10	138
HPV45			43	HPV45 E2	9	39
HPV45		10		HPV45 E2	10	39
HPV45		9	13 13	HPV45 E2	11	39
HPV45		10		HPV45 E2	8	326
HPV45		8 10	221	HPV45 E2	10	326
HPV45 HPV45		10	221 263	HPV45 E2	11	326
HPV45		11	263	HPV45 E2	10	98
HPV45		8	15	HPV45 E2	8	313
HPV45		9	215	HPV45 E2	11	313
HPV45		8	142	HPV45 E2	8	166
HPV45		10	142	HPV45 E2	11	166
HPV45		9	302	HPV45 E2	11	145
HPV45		8.	9	HPV45 E2	10	137
HPV45		9	9	HPV45 E2	11	137
HPV45		9	205	HPV45 E6	8	37
HPV45		10	205	HPV45 E6	11	59
HPV45		10	113	HPV45 E6	8	68
HPV45		11	113	HPV45 E6	11	68
HPV45		8	34	HPV45 E6	8	105
HPV45		9	34	HPV45 E6	11	105
HPV45		11	34	HPV45 E6	8	108
HPV45		8	229	HPV45 E6	8	18
HPV45		10	229	HPV45 E6	11	18
HPV45		9	208	HPV45 E6	8	32
HPV45		10	208	HPV45 E6	11	32
HPV45		10	276	HPV45 E6	8	16
HPV45		8	227	HPV45 E6	10	16
HPV45		10	227	HPV45 E6	10	51
HPV45		9	235	HPV45 E6	11	51
HPV45		11	235	HPV45 E6	8	143
HPV45		9	358	HPV45 E6	11	27
HPV45		10	358	HPV45 E6	9	20
HPV45		11	358	HPV45 E6	11	20
HPV45		10	155	HPV45 E6	9	77
HPV45		8	51	HPV45 E6	10	97
HPV45		10	51	HPV45 E6	9	88
HPV45		11	233	HPV45 E6	11	88
		~ *				

Table VIII HLA-A2 Supermotif-Bearing Peptides

		,	nLA-A2 Supermont-Bearing reputes	
HPV45 E6	10	43	HPV45 E7 8	83
HPV45 E6	8	47	HPV45 E7 10	83
HPV45 E6	9	47	HPV45 E7 8	41
HPV45 E6	8	53	HPV45 E7 10	41
HPV45 E6	9	53	HPV45 E7 8	20
HPV45 E6	11	53	HPV45 E7 8	74
HPV45 E6	10	136 .	HPV45 E7 11	74
HPV45 E6	9	132	HPV45 E7 8	91
HPV45 E6	11	120	HPV45 E7 8	97
HPV45 E6	8	30	HPV45 E7 9	44
HPV45 E6	9	30	HPV45 E7 9	47
HPV45 E6	10	30	HPV45 E7 8	14
HPV45 E6	11	130	HPV45 E7 11	14
HPV45 E6	10	60	HPV45 E7 11	11
HPV45 E6	9	93	HPV45 E7 8	В
HPV45 E6	10	93	HPV45 E7 8	87
HPV45 E6	11	93	HPV45 E7 9	87
HPV45 E6	8	54	HPV45 E7 10	75
HPV45 E6	10	54	HPV45 E7 8	17
HPV45 E6	11	54	HPV45 E7 10	17
HPV45 E6	9	36	HPV45 E7 11	17
HPV45 E6	10	92	HPV45 E7 9	57
HPV45 E6	11	92	HPV45 E7 10	23
HPV45 E6	9	13	HPV45 E7 10	89
HPV45 E6	11	13	HPV45 E7 8	88
HPV45 E6	11	102	HPV45 E7 11	88 54
HPV45 E6	9	25	HPV45 E7 9 HPV45 E7 10	54
HPV45 E6	8	1	HPV45 E7 10 HPV45 E7 8	5
HPV45 E6	8	95 · 95	HPV45 E7 9	5
HPV45 E6 HPV45 E6	9	22	HPV45 E7 11	5
HPV45 E6	10	22	HPV45 E7 10	72
HPV45 E6	8	114	HPV45 E7 8	85
HPV45 E6	11	111	HPV45 E7 10	85
HPV45 E6	8	7	HPV45 E7 11	85
HPV45 E6	11	7	HPV45 E7 8	80
HPV45 E6	8	149	HPV45 E7 11	80
HPV45 E6	10	149	HPV45 E7 11	93
HPV45 E6	11	146	HPV45 E7 9	7
HPV45 E6	8	41	HPV45 E7 9	86
HPV45 E6	9	29	HPV45 E7 10	86
HPV45 E6	10	29	HPV45 E7 10	94
HPV45 E6	11	29	HPV45 E7 11	94
HPV45 E6	8	24	HPV45 E7 9	76
HPV45 E6	10	24	HPV45 E7 11	76
HPV45 E6	10	84	HPV45 E7 10	12
HPV45 E6	11	84	HPV45 L1 11	517
HPV45 E6	8	89	HPV45 L1 10	
HPV45 E6	10	89	HPV45 L1 10	191
HPV45 E6	11	38	HPV45 L1 11	191
HPV45 E6	10	8	HPV45 L1 10	103
HPV45 E6	8	62	HPV45 L1 9	28
HPV45 E6	8	45	HPV45 L1 10	28
HPV45 E6	10	45	HPV45 L1 10	234
HPV45 E6	11	45	HPV45 L1 8	345
HPV45 E7	8	48	HPV45 L1 11	205
HPV45 E7	8	6	HPV45 L1 9	162
HPV45 E7	10	6	HPV45 L1 11 HPV45 L1 8	164 455
HPV45 E7	10	64	HPV45 L1 8 HPV45 L1 9	455
HPV45 E7	8	25	HPV45 LI 9	455

176

Table VIII HLA-A2 Supermotif-Bearing Peptides

HPV45	L1	В	374	HPV45	L1	11	351
HPV45	L1	11	374	HPV45	L1	9	10
HPV45	L1	8	184	HPV45	L1	11	10
HPV45	L1	9	184	HPV45	L1	8	503
HPV45	L1	9	276	HPV45	L1	11	143
HPV45	L1	11	252	HPV45	Ll	10	131
HPV45		9	409	HPV45	L1	8	137
HPV45		10	409	HPV45	L1	9	199
HPV45		10	188	HPV45	L1	9	306
HPV45		11	318	HPV45	L1	9	292
HPV45		10	480	HPV45		11	292
HPV45		В	401	HPV45	L1	8	435
HPV45		10	401	HPV45	L1	10	435
HPV45		11	401	HPV45		11	435
HPV45		9	301		L1	8	160
HPV45		11	301	HPV45	L1	11	160
HPV45		9	226	HPV45	L1	8	62
HPV45		10	226		L1	9	62
HPV45		10	490		L1	10	62
HPV45		11	490	HPV45		10	396
HPV45		8	155	HPV45		8	221
HPV45		9	155		L1	10	221
HPV45		10	155	HPV45		9	12
HPV45		11	155	HPV45	L1	8	11
HPV45		11	229	HPV45	L1	10	11
HPV45		8	242	HPV45		8	5
HPV45		10	242	HPV45		9	5
HPV45		8	364	HPV45	L1	11	5
HPV45		9	364	HPV45	L1	9	428
HPV45		10	364	HPV45	L1	8	185
HPV45		9	296	HPV45	L1	8	411
HPV45	L1	9	446	HPV45	L1	11	411
HPV45	L1	10	446	HPV45	L1	9	166
HPV45	L1	11	446	HPV45	L1	11	166
HPV45	L1	8	169	HPV45	L1	8	328
HPV45	L1	8	133	HPV45	Ll	9	344
HPV45	L1	10	133	HPV45	L1	8	152
HPV45	L1	8	121	HPV45	L1	10	152
HPV45	L1	10	121	HPV45	L1	11	152
HPV45	L1	9	416	HPV45		9	473
HPV45	L1	10	246	HPV45	L1	9	86
HPV45	L1	9	283	HPV45		8	467
HPV45	L1	11	283	HPV45	L1	8	179
HPV45		8	404		L1	11	179
HPV45	Ll	9	404	HPV45		9	91
HPV45	L1	10	404		L1	11	68
HPV45	L1	11	404	HPV45		8	240
HPV45		10	14	HPV45	L1	10	240
HPV45		8	406	HPV45		9	402
HPV45		9	406	HPV45	L1	10	402
HPV45		10	406		L1	11	402
HPV45		8	450	HPV45	L1	9	207
HPV45		8	359	HPV45		11	207
HPV45		11	359	HPV45		9	413
HPV45		8	82	HPV45	L1	11	371
HPV45		11	82	HPV45	L1	10	69
HPV45		8	233	HPV45	L1	11	69
HPV45		11	233	HPV45		8	444
HPV45		8	351	HPV45	L1	11	444
HPV45	T.1	10	351	HPV45	r1	11	499

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-AZ Supermotif-Bearing Peptides		
HPV45 L1	8	125	HPV45 L1	9	290
HPV45 L1	10	125	HPV45 L1	11	290
HPV45 L1	11	1	HPV45 L1	9	124
HPV45 L1	10	27	HPV45 L1	11	124
HPV45 L1	11	27	HPV45 L1	8	56
HPV45 L1	8	227	HPV45 L1	9	46
HPV45 L1	9	227	HPV45 L1	9	265
HPV45 L1	8	4	HPV45 L1	8	158
HPV45 L1	9	4	HPV45 L1	9	158
HPV45 L1	10	4	HPV45 L1	10	158
HPV45 L1	8	370	HPV45 L1	10	93
HPV45 L1	10	310	HPV45 L1	11	93
HPV45 L1	8	356	HPV45 L1	9	254
HPV45 L1	10	356	HPV45 L1	11	254
HPV45 L1	11	356	HPV45 L1	11	58
HPV45 L1	9	49	HPV45 L1	10	427
HPV45 L1	11 9	49	HPV45 L1	9	327
HPV45 L1		219	HPV45 L1 HPV45 L1	9	443 272
HPV45 L1 HPV45 L1	10 9	219 19	HPV45 L1 HPV45 L1	11	333
HPV45 L1	10	19	HPV45 L1	10	521
HPV45 L1	11	19	HPV45 L1	8	115
HPV45 L1	9	17	HPV45 L1	11	115
HPV45 L1	11	17	HPV45 L1	10	238
HPV45 L1	9	173	HPV45 L1	8	368
HPV45 L1	11	173	HPV45 L1	10	368
HPV45 L1	8	516	HPV45 L1	9	376
HPV45 L1	8	190	HPV45 L1	9	519
HPV45 L1	11	190	HPV45 L1	10	35
HPV45 L1	8	22	HPV45 L1	11	35
HPV45 L1	8	248	HPV45 L1	8	43
HPV45 L1	8	214	HPV45 L1	10	453
HPV45 L1	9	214	HPV45 L1	11	453
HPV45 L1	8	493	HPV45 L1	9	175
HPV45 L1	10	493	HPV45 L1	11	175
HPV45 L1	9	299	HPV45 L1 HPV45 L1	8	414 414
HPV45 L1	11	299 508	HPV45 L1 HPV45 L1	11 9	522
HPV45 L1 HPV45 L1	10	508	HPV45 L1	11	522
HPV45 L1	11 11	387	HPV45 L1	8	163
HPV45 L1	8	440	HPV45 L1	9	509
HPV45 L1	9	380	HPV45 L1	10	509
HPV45 L1	9	501	HPV45 L1	8	220
HPV45 L1	10	501	HPV45 L1	9	220
HPV45 L1	8	87	HPV45 L1	11	220
HPV45 L1	8	182	HPV45 L1	8	410
HPV45 L1	10	182	HPV45 L1	9	410
HPV45 L1	11	182	HPV45 L1	10	116
HPV45 L1	8	407	HPV45 L1	11	116
HPV45 L1	9	407	HPV45 L1	10	372
HPV45 L1	11	407	HPV45 L1	8	200
HPV45 L1	11	281	HPV45 L1	9	239
HPV45 L1	9	334	HPV45 L1 HPV45 L1	11	239 412
HPV45 L1 HPV45 L1	9	357 357	HPV45 L1 HPV45 L1	8	167
HPV45 L1	10 10	206	HPV45 L1	10	167
HPV45 L1	11	26	HPV45 L1	9	181
HPV45 L1	11	263	HPV45 L1	11	181
HPV45 L1	10	180	HPV45 L1	8	377
HPV45 L1	8	290	HPV45 L1	9	122
	-				

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			TILA-A2 Supermon-Bear	ng repudes		
HPV45	L1	11	122	HPV45 L2	9	278
HPV45	L1	8	365	HPV45 L2	10	278
HPV45		9	365	HPV45 L2	8	322
HPV45	1.1	11	365	HPV45 L2	11	322
HPV45		11	441	HPV45 L2	9	142
HPV45		9	70	HPV45 L2	11	142
HPV45		10	70	HPV45 L2	11	16
HPV45		8	297	HPV45 L2	9	260
HPV45		11	297	HPV45 L2	9	83
HPV45		9	361	HPV45 L2	10	83
HPV45		11	361	HPV45 L2	11	83
HPV45		9	36	HPV45 L2	9	30
HPV45		10	36	HPV45 L2	10	30 .
HPV45		11	102	HPV45 L2	11	30 .
HPV45		11	44	HPV45 L2	10	397
HPV45		9	454	HPV45 L2	11	397
HPV45		10	454	HPV45 L2	10	348
HPV45		10	165	HPV45 L2	8	331
HPV45		8	293	HPV45 L2	10	331
HPV45		10	293	HPV45 L2	11	331
HPV45		8	417	HPV45 L2	8	194
HPV45		10	500	HPV45 L2	8	129
HPV45		11	500	HPV45 L2	9	129
HPV45			456	HPV45 L2	11	129
HPV45		8 10	360	HPV45 L2	8	333
HPV45		8	362	HPV45 L2	9	333
HPV45		10	362	HPV45 L2	8	169
				HPV45 L2		175
HPV45		11	362 47	HPV45 L2	8 10	175
HPV45		8		HPV45 L2		456
		11	47 78	HPV45 L2	8	200
HPV45		11		HPV45 L2	11	200
HPV45		8	127	HPV45 L2	9	53
HPV45		10	196	HPV45 L2	8	241
HPV45		9	420	HPV45 L2		
HPV45		10	420		9	241
HPV45		9	303	HPV45 L2	1.0	241
HPV45		8	38	HPV45 L2 HPV45 L2	11	276
HPV45		10	38		9	122
HPV45		11	38	HPV45 L2	10	122
HPV45		8	95	HPV45 L2	11	157
HPV45		9	95	HPV45 L2	8	306
HPV45		10	95	HPV45 L2	8	368
HPV45		11	53	HPV45 L2	9	368
HPV45		9	6	HPV45 L2	10	368
HPV45		10	6	HPV45 L2	8	116
HPV45		8	381	HPV45 L2	10	116
HPV45		9	381	HPV45 L2	9	51
HPV45		10	381	HPV45 L2	11	51
HPV45		8	327	HPV45 L2	9	430
HPV45		11	327	HPV45 L2	8	300
HPV45		8	286	HPV45 L2	8	25
HPV45		9	286	HPV45 L2	11	25
HPV45		10	328	HPV45 L2	8	206
HPV45		11	328	HPV45 L2	10	206
HPV45		11	303	HPV45 L2	10	60
HPV45		10	340	HPV45 L2	8	124
HPV45		8	139	HPV45 L2	8	37
HPV45		9	139	HPV45 L2	9	37
HPV45		8	405	HPV45 L2	8	134
HPV45	L2	10	405	HPV45 L2	10	134

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			nLA-A2 Supermout-Beatt	ng repud	C3		
HPV45	L2	11	134	HPV45	L2	8	325
HPV45	L2	8	292	HPV45	L2	10	325
HPV45	L2	8	411	HPV45	L2	9	209
HPV45		11	411	HPV45		10	209
HPV45		9	326	HPV45		8	399
HPV45		10	167	HPV45		9	399
HPV45		9	406	HPV45		10	399
HPV45		8	279	HPV45		11	258
HPV45		9	279	HPV45		8	73
EPV45		8	407	HPV45		9	73
HPV45	L2	9	44	HPV45	I.2	10	73
HPV45		11	44	HPV45		11	336
HPV45		8	143	HPV45		8	214
HPV45		10	143	HPV45		8	391
HPV45		8	130	HPV45	L2	10	391
HPV45		10	130	HPV45		9	413
HPV45		11	130	HPV45		11	413
HPV45		10	103	HPV45	L2	10	171
HPV45	1.2	11	103	HPV45	L2	8	98
HPV45		8	43	HPV45		9	98
HPV45		10	43	HPV45	L2	10	98
HPV45	L2	10	22	HPV45	L2	11	120
HPV45	L2	11	22	HPV45	L2	8	420
HPV45		8	34	HPV45	L2	10	420
HPV45		11	34	HPV45	L2	8	86
HPV45	L2	11	40	HPV45	L2	11	86
HPV45	L2	10	337	HPV45	L2	11	185
HPV45	L2	8	334	HPV45	L2	11	257
HPV45	L2	11	197	HPV45	L2	8	145
HPV45	L2	8	45	HPV45	L2	11	145
HPV45	L2	10	45	HPV45	L2	11	216
HPV45	L2	8	242	HPV45	L2	9	95
HPV45	L2	9	242	HPV45	L2	10	95
HPV45	L2	8	375	HPV45	L2	11	95
HPV45	L2	9	392	HPV45	L2	8	118
HPV45	L2	11	392	HPV45	L2	10	90
HPV45	L2	8	106	HPV45	L2	11	232
HPV45	L2	9	106	HPV45	L2	10	198
HPV45	L2	8	248	HPV45	L2	9	172
HPV45	L2	8	422	HPV45	L2	11	172
HPV45	L2	10	422	HPV45	L2	10	233
HPV45	L2	8	179	HPV45	L2	11	233
HPV45	L2	8	231	HPV45	L2	8	5
HPV45	L2	9	79	HPV45	L2	10	5
HPV45	L2	8	270	HPV45		11	5
HPV45	L2	10	270	HPV45		8	229
HPV45		11	270	HPV45		10	229
HPV45	L2	9	387	HPV45	L2	10	11
HPV45	L2	8	160	HPV45	L2	11	451
HPV45	L2	9	160	HPV45	L2	8	298
HPV45		11	160	HPV45		10	298
HPV45	L2	9	285	HPV45		10	225
HPV45	L2	10	285	HPV45		8	19
HPV45	L2	8	356	HPV45	L2	8	316
HPV45	L2	9	356	HPV45		11	316
HPV45	L2	9	138	HPV45		11	220
HPV45	L2	10	138	HPV45		8	235
HPV45	L2	8	254	HPV45		9	235
HPV45	L2	9	254	HPV45		8	13
HPV45	L2	10	254	HPV45	L2	8	339

Table VIII HLA-A2 Supermotif-Bearing Peptides

			IILA-7	12 Supermont-Deating i	reptide	23		
HPV45	L2	11	339	H	PV45	L2	11	249
HPV45	L2	9	394	HI	PV45	L2	9	104
HPV45		8	166		PV45		10	104
HPV45		11	166		PV45		11	104
HPV45		8	151		PV45		8	388
HPV45		11	151		PV45		11	388
HPV45		11	102		PV45		В	112
HPV45		9	49		PV45		10	112
HPV45		11	49		PV45		11	81
HPV45		8	374		PV45		9	91
HPV45		9	374		PV45		8	350
HPV45		9	247		PV45		10	350
HPV45		10	239		PV45		11	350
HPV45		11	239		PV45		11	428
HPV45		10	379		PV45		8	401
HPV45		11	379		PV45		10	71
HPV45		8	362		PV45		11	71
HPV45		10	212		PV56		8	15
HPV45		8	154		PV56		11	15
HPV45		9	417		PV56		8	21
HPV45		10	417		PV56		9	21
HPV45		11	417		PV56		9	4
HPV45		8	424		PV56		10	71
HPV45		11	424		PV56		8	204
HPV45		10	149		PV56		11	204
HPV45		9	111		PV56		8	39
HPV45		11	111		PV56		9	39
HPV45		9	380		PV56		9	263
HPV45		10	380		PV56		11	263
HPV45		11	380		PV56		10	117
HPV45		10	262		PV56		8	288
HPV45		8	105		PV56		11	288
HPV45		9	105		PV56		8	154
HPV45		10	105		PV56		11	154
HPV45		10	304		PV56		9	128
HPV45		8	38		PV56		9	17
HPV45		8	261		PV56		11	17
HPV45		11	261		PV56		9	294
HPV45		8	136		PV56		10	294
HPV45		9	136		PV56		9	254
HPV45		11	136		PV56		11	254
HPV45		11	359		PV56		11	261
HPV45		9	135		PV56		11	99
HPV45		10	135		PV56		10	94
HPV45		10	425		PV56		9	201
HPV45		9	426		PV56		11	201
HPV45		10	146		PV56		8	210
HPV45		10	389		PV56		9	239
HPV45		11	293		PV56		10	208
HPV45		10	217		PV56		10	297
HPV45		8	80		PV56		11	297
HPV45					PV56		8	20
HPV45		9 11	113 113		PV56		9	20
HPV45		8	92		PV56		10	20
HPV45		8	31		PV56		10	280
					PV56		9	280
HPV45		9	31		PV56			11
HPV45		10	31		PV56		8	
HPV45		11	31		PV56		10	11
HPV45		8	140		PV56		11	11
HPV45	ьz	11	140	H	2 42 0	54	8	9

Table VIII HLA-A2 Supermotif-Bearing Peptides

			TILA-A2 Supermont-Beam	ng repno	CS		
HPV56	E2	10	9	HPV56	E2	8	264
HPV56	E2	8	299	HPV56	E2	10	264
HPV56		9	299	HPV56	E2	10	205
HPV56	E2	8	258	HPV56	E2	11	237
HPV56	E2	10	258	HPV56	E2	8	88
HPV56	E2	8	233	HPV56	E2	10	35
HPV56	E2	8	163	HPV56	E2	11	270
HPV56	E2	10	163	HPV56	E2	8	111
HPV56	E2	11	108	HPV56	E2	8	102
HPV56	E2	11	90	HPV56	E2	11	102
HPV56	E2	8	5	HPV56	E6	11	89
HPV56	E2	11	5	HPV56	E6	8	64
HPV56	E2	9.	72	HPV56	E6	8	139
HPV56	E2	10	1	HPV56	E6	8	69
HPV56	E2	9	216	HPV56	E6	8	33
HPV56	E2	8	160	HPV56	E6	9	33
HPV56	E2	9	160	HPV56	E6	11	33
HPV56	E2	11	160	HPV56	E6	9	23
HPV56		8	149	HPV56	E6	11	39
HPV56		10	152	HPV56		8	20
HPV56		9	19	HPV56		10	20
HPV56		10	19	HPV56		8	44
HPV56		11	19	HPV56	E6	10	44
HPV56		10	6	HPV56		8	48
HPV56		11	6	HPV56	E6	9	48
HPV56		8	14	HPV56	E6	8	88
HPV56		9	14	HPV56	E6	8	129
HPV56		11	279	HPV56	E6	8	17
HPV56		8	135	HPV56 HPV56		10 11	17 17
HPV56		11	135 144	HPV56		10	131
HPV56 HPV56		10 9	272	HPV56	E6	9	94
HPV56		10	272		E6	10	94
HPV56		11	272	HPV56		11	94
HPV56		9	169	HPV56		11	54
HPV56		10	169	HPV56		8	97
HPV56		11	169	HPV56	E6	11	130
HPV56		11	26	HPV56	E6	9	26
HPV56		8	266	HPV56	E6	11	103
HPV56		8	171	HPV56	E6	8	113
HPV56	E2	9	171	HPV56	E6	10	40
HPV56	E2	10	141	HPV56	E6	10	55
HPV56	E2	8	282	HPV56	E6	11	55
HPV56	E2	9	28	HPV56		10	25
HPV56		10	28	HPV56		9	112
HPV56		9	259		E6	8	8
HPV56		9	36	HPV56		11	8
HPV56		11	36	HPV56		10	145
HPV56		10	289	HPV56		11	145
HPV56		9	206	HPV56		8	42
HPV56		10	27	HPV56		10	42 30
HPV56		11	27 167	HPV56 HPV56		11	144
HPV56 HPV56		11 8	167	HPV56		10	67
HPV56		9	164	HPV56	E6	10	93
HPV56		10	155		E6	11	93
HPV56		11	155	HPV56		8	14
HPV56		8	18	HPV56	E6	9	14
HPV56		10	18	HPV56	E6	11	14
HPV56		11	18	HPV56		10	85

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-	A2 Supermotit-Bearing	Peptide	es		
HPV56	E6	11	85	H:	PV56	E7	11	51
HPV56	E6	10	90	H	PV56	E7	8	84
HPV56	E6	9	21	H	PV56	E7	10	84
HPV56	E6	11	21	H	PV56	E7	11	84
HPV56	E6	8	63	H	PV56	E7	8	78
HPV56	E6	9	63	H:	PV56	E7	9	78
HPV56	È7	8	93			E7	9 .	7
HPV56		10	93		PV56		10	7
HPV56		9	75			E7	8	95
HPV56		11	75			E7	8	12
HPV56		8	22			E7	10	12
HPV56		8	82		PV56		8	72
HPV56		9	82		PV56		8	86
HPV56		10	82		PV56 PV56		9	86
HPV56		10	10 20		PV56		10 11	86 86
HPV56		8	14		PV56		9	11
HPV56		10	14		PV56		9 11	11
HPV56		10	70		PV56		9	71
HPV56		9	92		PV56		9	85
HPV56		11	92				10	85
HPV56		9	42		PV56		11	85
HPV56		8	56		PV56		11	458
HPV56		10	62		PV56		10	198
HPV56		11	62		PV56		11	198
HPV56		8	76		PV56			350
HPV56		10	76		PV56		10	338
HPV56	E7	11	76	H	PV56	L1	9	58
HPV56	E7	8	54	HI	PV56	L1	11 -	58
HPV56	E7	10	54	HI	PV56	L1	10	381
HPV56	E7	8	4	H	PV56	L1	8	327
HPV56	E7	9	4	H	PV56	L1	9	327
HPV56		10	4				8	514
HPV56		8	89		PV56		10	514
HPV56		9	89		PV56		11	444
HPV56		8	90		PV56		10	37
HPV56		11	90		PV56			512
HPV56		8	8					207
HPV56		9	8		PV56			207
HPV56		8 11	43		PV56 PV56		10 9	207 79
HPV56		9	15		PV56			79
HPV56		9	94					26
HPV56		11	47		PV56		-	26
HPV56		8	6		PV56		8	19
HPV56		10	6		PV56		9	19
HPV56		11	6		PV56		11	19
HPV56		9	52		PV56			191
HPV56	E7	10	52	H	PV56	L1	9	191
HPV56	E7	9	49	H	PV56	L1	8	461
HPV56	E7	11	73	H	PV56	L1	10	195
HPV56		8	88	H	PV56	Ь1	9	389
HPV56	E7	9	88	H	PV56	L1	11	274
HPV56	E7	10	88		PV56			233
HPV56		10	48					233
HPV56		8	87		PV56		8	128
HPV56		9	87		PV56		10	128
HPV56		10	87					493
HPV56		11	87		PV56		11	493
HPV56	E7	10	51	H	PV56	r1	8	162

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotit-Beari	ng Peptides		
HPV56	L1	11	236	HPV56 L1	8	413
HPV56	L1	8	369	HPV56 L1	9	413
HPV56		9	369	HPV56 L1	10	413
HPV56	L1	10	369	HPV56 L1	11	270
HPV56		10	23	HPV56 L1	8	186
HPV56	L1	10	481	HPV56 L1	11	186
HPV56	L1	11	337	HPV56 L1	9	93
HPV56	L1	8	404	HPV56 L1	8	300
HPV56		11	404	HPV56 L1	10	300
HPV56		8	383	HPV56 L1	8	245
HPV56	L1	11	383	HPV56 L1	10	245
HPV56	Li	11	464	HPV56 L1	9	98
HPV56	L1	9	303	HPV56 L1	9	55
HPV56	L1	8	140	HPV56 L1	9	45
HPV56	L1	10	140	HPV56 L1	9	474
HPV56	L1	9	419	HPV56 L1	11	474
HPV56	L1	11	419	HPV56 L1	8	222
HPV56	L1	10	253	HPV56 L1	10	78
HPV56	L1	9	290	HPV56 L1	11	78
HPV56	L1	9	21	HPV56 L1	11	224
HPV56	L1	8	409	HPV56 L1	11	77
HPV56	L1	10	409	HPV56 L1	9	431
HPV56	L1	8	407	HPV56 L1	11	502
HPV56	L1	9	407	HPV56 L1	9	484
HPV56	L1	10	407	HPV56 L1	8	247
HPV56	L1	8	364	HPV56 L1	10	247
HPV56	L1	11	364	HPV56 L1	10	405
HPV56	L1	9	148	HPV56 L1	11	405
HPV56	L1	8	240	HPV56 L1	11	347
HPV56		9	206	HPV56 L1	8	132
HPV56	L1	10	206	HPV56 L1	10	132
HPV56		11	206	HPV56 L1	11	36
HPV56		8	25	HPV56 L1	9	421
HPV56		10	25	HPV56 L1	11	421
HPV56		8	356	HPV56 L1	8	234
HPV56		10	356	HPV56 L1	9	234
HPV56		11	356	HPV56 L1	8	333
HPV56		8	17	HPV56 L1	8	1
HPV56		10	17	HPV56 L1 HPV56 L1	10	5 503
HPV56		11	17			
HPV56		10	138	HPV56 L1 HPV56 L1	11	503 376
HPV56		11	150 438	HPV56 L1	10	428
HPV56		10	438	HPV56 L1	8	436
HPV56 HPV56			144	HPV56 L1	9	436
HPV56		8	506	HPV56 L1	10	436
HPV56		9	506	HPV56 L1	9	180
HPV56		10	506	HPV56 L1	11	180
HPV56		8	71	HPV56 L1	10	123
HPV56		9	71	HPV56 L1	11	123
HPV56		10	71	HPV56 L1	8	167
HPV56		10	399	HPV56 L1	8	430
HPV56		10	459	HPV56 L1	10	430
HPV56		8	414	HPV56 L1	8	483
HPV56		9	414	HPV56 L1	10	483
HPV56		11	414	HPV56 L1	8	375
HPV56		8	192	HPV56 L1	11	375
HPV56		8	251	HPV56 L1	8	361
HPV56		9	251	HPV56 L1	10	361
HPV56		9	392	HPV56 L1	11	361
		-				

Table VIII HLA-A2 Supermotif-Bearing Peptides

				ILA-AZ Supermoni-nea	ing repud	es		
HPV56	Ll	9	91		HPV56	L1	8	520
HPV56	L1	11	91		HPV56	L1	10	520
HPV56	L1	9	226		HPV56	L1	10	100
HPV56	L1	10	226		HPV56	Ll	11	67
HPV56	L1	9	28		HPV56	L1	9	446
HPV56	L1	10	28		HPV56	L1	9	332
HPV56	Ll	11	28		HPV56	L1	11	279
HPV56	L1	10	172		HPV56	L1	9	261
HPV56	L1	8	197		HPV56	L1	11	261
HPV56	L1	11	197		HPV56	Ll	9	489
HPV56	L1	11	511		HPV56	L1	8	373
	L1	8	228		HPV56	Ll	9	373
HPV56		10	228		HPV56	L1	10	373
HPV56		8	31		HPV56	L1	9	182
HPV56		10	473		HPV56	Ll	11	182
HPV56		9	221		HPV56	L1	9	86
HPV56		8	255		HPV56	L1	11	86
	L1	9	155		HPV56	Ll	11	323
HPV56		11	146		HPV56	L1	11	380
HPV56		8	496		HPV56	L1	8	304
HPV56		10	496		HPV56	L1	9	377
	L1	8	13		HPV56	Li	8	415
	L1	10	13		HPV56 HPV56	L1	10	188
HPV56		11	4		HPV56		9 11	
HPV56		8	467		HPV56	L1 L1	11	188 212
HPV56 HPV56	L1	9	467 50		HPV56	L1	8	215
	L1	9	50		HPV56	L1	9	215
HPV56		10	50	_		L1	10	215
	L1	8	522		HPV56	Li	11	215
HPV56		8	522		HPV56	Li	8	370
	Li	8	189		HPV56	LI	9	370
	Li	10	189		HPV56	L1	11	370
HPV56		11	189		HPV56	Li	9	366
	Li	9	410		HPV56	Li	11	366
	L1	11	410		HPV56	L1	10	57
	L1	11	288		HPV56	L1	8	326
HPV56		9	339		HPV56	L1	9	326
	L1	9	362		HPV56	L1	10	326
HPV56	Ll	10	362		HPV56	L1	9	513
HPV56	L1	9	504		HPV56	L1	11	513
HPV56	L1	10	504		HPV56	L1	9	173
HPV56	L1	11	504		HPV56	L1	11	173
HPV56	L1	10	384		HPV56	L1	9	246
	L1	10	187		HPV56	L1	11	246
HPV56	L1	10	213		HPV56	L1	8	420
HPV56		11	213		HPV56	L1	10	420
HPV56		8	297		HPV56		11	259
HPV56		9	297		HPV56	L1	8	87
	L1	11	297		HPV56	L1	10	87
	L1	9	349		HPV56	L1	9	214
HPV56		10	159		HPV56	L1	10	214
HPV56		11	159		HPV56	L1	11	214
HPV56		10	110		HPV56	L1	10	365
HPV56		9	131		HPV56	L1	8	56
HPV56		11	131		HPV56	L1	11	56
HPV56		8	65		HPV56	ы	8	367
HPV56		9	272		HPV56	L1	10	367
HPV56		8	417		HPV56	ь1	11	367
HPV56	L1	11	417		HPV56	L1	8	134

Table VIII
HLA-A2 Supermotif-Bearing Peptides

			HLA-A2 Supermotif-Bearing Peptides		
HPV56 L1	10	203	HPV56 L2	11	398
HPV56 L1	8	7	HPV56 L2	8	175
HPV56 L1	9	423	HPV56 L2	10	175
HPV56 L1	10	423	HPV56 L2	8	457
HPV56 L1	8	268	HPV56 L2	8	382
HPV56 L1	10	47	HPV56 L2	9	382
HPV56 L1	11	47	HPV56 L2	10	382
HPV56 L1	10	396	HPV56 L2	8	200
HPV56 L1	9	283	HPV56 L2	11	200
HPV56 L1	8	102	HPV56 L2	9	162
HPV56 L1	10	102	HPV56 L2	8	241
HPV56 L1	9	325	HPV56 L2	9	241
HPV56 L1	10	325	HPV56 L2	11	241
HPV56 L1	11	325	HPV56 L2	11	276
HPV56 L1	11	62	HPV56 L2	10	231
HPV56 L1	8	453	HPV56 L2	9	122
HPV56 L2	9	222	HPV56 L2	10	122
HPV56 L2	8	281	HPV56 L2	8	287
HPV56 L2	9	281	HPV56 L2	9	51
HPV56 L2	8	327	HPV56 L2	11	51
HPV56 L2	11	327	HPV56 L2	9	418
HPV56 L2	9	303	HPV56 L2	10	418
HPV56 L2	11	303	HPV56 L2	8	116
HPV56 L2	10	246	HPV56 L2	10	314
HPV56 L2	9	367	HPV56 L2	8	188
HPV56 L2	10	14	HPV56 L2	10	188
HPV56 L2	9	6	HPV56 L2	8	56
HPV56 L2	10	6	HPV56 L2	8	360
HPV56 L2 ·	10	201	HPV56 L2	9	360
HPV56 L2	8	139	HPV56 L2	9	346
HPV56 L2	11	139	HPV56 L2	8	25
HPV56 L2	8	322	HPV56 L2	11	25
HPV56 L2	11	322	HPV56 L2	8	206
HPV56 L2	8	142	HPV56 L2	10	206
HPV56 L2	9	142	HPV56 L2	8	62
HPV56 L2	11	142	HPV56 L2	11	62
HPV56 L2	10	406	HPV56 L2	10	60
HPV56 L2	10	349	HPV56 L2	8	310
HPV56 L2	11	260	HPV56 L2	9	269
HPV56 L2	8	425	HPV56 L2	11	269
HPV56 L2	9	83	HPV56 L2	11	293
HPV56 L2	10	83	HPV56 L2	8	156
HPV56 L2	11	83	HPV56 L2	8	372
HPV56 L2	10	30	HPV56 L2 HPV56 L2	9	372 151
HPV56 L2	10	429 429	HPV56 L2 HPV56 L2	11	151
HPV56 L2 HPV56 L2	11	357	HPV56 L2	10	221
HPV56 L2	10 11	357	HPV56 L2	9	326
HPV56 L2	8	169	HPV56 L2	10	180
HPV56 L2	8	331	HPV56 L2	9	44
HPV56 L2	10	331	HPV56 L2	8	432
HPV56 L2	8	194	HPV56 L2	11	432
HPV56 L2	9	194	HPV56 L2	9	305
HPV56 L2	8	129	HPV56 L2	11	305
HPV56 L2	9	129	HPV56 L2	11	157
HPV56 L2	11	129	HPV56 L2	8	143
HPV56 L2	8	333	HPV56 L2	10	143
HPV56 L2	9	36	HPV56 L2	8	130
HPV56 L2	10	36	HPV56 L2	10	130
HPV56 L2	10	398	HPV56 L2	11	130

186

Table VIII HLA-A2 Supermotif-Bearing Peptides

HPV56	L2	8	279	HPV56 L2	10	73
HPV56		10	279	HPV56 L2	8	420
HPV56		11 '	279	HPV56 L2	11	420
HPV56		11	81	HPV56 L2	10	171
HPV56	L2	9	407	HPV56 L2	11	171
HPV56	L2	10	103	HPV56 L2	8	98
HPV56	L2	11	103	HPV56 L2	9	98
HPV56	L2	9	91	HPV56 L2	10	98
HPV56	L2	8	229	HPV56 L2	10	410
HPV56	L2	9	229	HPV56 L2	11	185
HPV56	L2	10	302	HPV56 L2	8	145
HPV56	L2	8	34	HPV56 L2	11	145
HPV56	L2	11	34	HPV56 L2	8	166
HPV56	L2	8	43	HPV56 L2	11	166
HPV56	L2	10	43	HPV56 L2	10	328
HPV56	L2	10	22	HPV56 L2	11	328
HPV56		11	22	HPV56 L2	8	16
HPV56		8	19	HPV56 L2	11	16
HPV56		8	38	HPV56 L2	9	232
HPV56		8	235	HPV56 L2	11	232
HPV56		8	263	HPV56 L2	10	198
HPV56		9	263	HPV56 L2	9	172
HPV56		9	181	HPV56 L2	10	172
HPV56		11	337	HPV56 L2	11	172
HPV56		8	45	HPV56 L2	8	306
HPV56		8	106	HPV56 L2	10	306
HPV56		8	248	HPV56 L2	8	233
HPV56		11	197	HPV56 L2	10	233
HPV56			353	HPV56 L2	10	11
HPV56		9	179	HPV56 L2	8	5
HPV56		8 11	179	HPV56 L2	10	5
			278	HPV56 L2	11	5
HPV56		9		HPV56 L2		220
HPV56		11	278	HPV56 L2	11	
HPV56		9	385	HPV56 L2	11	452 298
HPV56		9	388	HPV56 L2	8 10	298
HPV56		8	239	HPV56 L2		225
HPV56		9	239	HPV56 L2	10 8	316
HPV56		10	239	HPV56 L2	11	316
HPV56		11	239			
HPV56		10	285	HPV56 L2	10	250
HPV56		8	8.6	HPV56 L2	10	370
HPV56		11	86	HPV56 L2	-11	370
H₽V56		11	245	HPV56 L2	9	339
HPV56		9	138	HPV56 L2	8	13
HPV56		8	325	HPV56 L2	11	13
H₽V56		10	325	HPV56 L2	11	102
HPV56		10	374	HPV56 L2	9	262
HPV56		8	214	HPV56 L2	10	262
HPV56		9	214	HPV56 L2	9	49
HPV56		10	214	HPV56 L2	11	49
HPV56		10	90	HPV56 L2	9	363
HPV56		8	254	HPV56 L2	11	363
HPV56		10	254	HPV56 L2	8	154
HPV56		11	254	HPV56 L2	10	154
HPV56	L2	8	160	HPV56 L2	9	79
HPV56	L2	9	160	HPV56 L2	9	378
HPV56	L2	11	160	HPV56 L2	10	378
HPV56	L2	8	392	HPV56 L2	10	212
HPV56		8	73	HPV56 L2	11	212
HPV56	L2	9	73	HPV56 L2.	8	134

Table VIII HLA-A2 Supermotif-Bearing Peptides

				IILA-AZ SI	ірегиюш-всаги	ig repud	es		
HPV56	L2	10	134			HPV56	L2	10	71
HPV56	L2	11	134			HPV56	L2	11	71
HPV56	L2	8	148						
HPV56	L2	10	148						
HPV56	L2	11	148						
HPV56	L2	9	365						
HPV56	L2	11	365						
HPV56		9	95						
HPV56	L2	10	95						
HPV56		11	95						
HPV56		9	111						
HPV56		10	390						
HPV56		8	304						
HPV56		10	304						
HPV56		8	80						
HPV56		8	379						
HPV56		9	379						
HPV56		11	379						
HPV56		9	105						
HPV56 HPV56		9	105 247						
HPV56		9	15						
HPV56		8	386						
HPV56		11	386						
HPV56		8	136						
HPV56		9	136						
HPV56		11	136						
HPV56		9	135						
HPV56		10	135						
HPV56		9	149						
HPV56	L2	10	149						
HPV56	L2	11	2						
HPV56	L2	9	280						
HPV56	L2	10	280						
HPV56		8	270						
HPV56		10	270						
HPV56		11	270						
HPV56		8	366						
HPV56		10	366						
HPV56		10	167						
HPV56		8	112						
HPV56		10	140						
HPV56 HPV56		11	140 408						
HPV56		8	389						
HPV56		11	389						
HPV56		11	236						
HPV56		9	104						
HPV56		10	104						
HPV56		8	84						
HPV56		9	84						
HPV56		10	84						
HPV56		8	92						
HPV56		9	31						
HPV56		11	31						
HPV56		11	40						
HPV56	L2	8	351						
HPV56	L2	11	351						
HPV56	L2	8	431						
HPV56	L2	9	431						

188

		11011110	supermont a upride.		
2	3	4	E1	10	407
Ll	9	234	E4	8	61
L2	10	329	E4	9	61
L2	11	329	L2	10	14
E5	8	9	L2	11	14
E5	9	9	E1	9	525
E5	10	9	E1	11	525
E1	8	318	E6	10	10
E1	10	318	E6	8	86
L1	8	489	E1	11	77
L1	10	489	E1	10	101
L2	9	340	L1	9	43
L2	11	340	E2	1,0	231
E4	8	2	L1	8	483
E4	10	2	E1	8	601
E2	8	3	E6	11	64
L2	8 9	286 286	E1 E2	11	234
L2 E2	8	72	E2	9	124
E2	11	72	L1	10 9	124 341
L2	10	112	Li	11	341
L2	11	112	E1	11	406
E1	11	112	E1	10	473
L2	8	140	E6	8	67
L2	11	140	E6	9	137
Ll	8	420	E2	9	296
El	8	475	E2	8	35
E1	9	22	E2	9	35
E1	11	22	E1	8	488
E2	10	250	E1	11	488
E1	8	65	L1	9	153
El	10	65	E2	8	11
E4	9	14	E7	9	71
E4	11	14	E1	9	14
L2	8	228	E1	10	14
L2	11	228	E1	11	14
L1	11	81	L1	8	171
L2	9	421	E6	8	131
L2	10	421	E2	8	252
E1	10	554	E6	8	31
E1	11	554	E6	9	31
E6	11	37	E6	10	31
E5	8	79	E6	11	31
E5	9	79	E1	9	640
E5	11	79	E1	10	640
E1	9	319	E4	10	64
L1	9	203	E4	11	64
L2	9	327	E2	8	9
E1	10	63	E2	10	9
L2	8	341	E2	8	153 516
L2	10	341	E1 E1	10 11	516
L1 L1	8 10	312 300	E1	8	524
E4	9	3 0 0	E1	10	524
E4	11	3	E2	11	230
E1	10	381	L1	9	24
E2	11	217	E1	8	369
L1	9	217	E1	9	369
L1	11	22	E1	8	170
E1	11	296 .	E1	10	170
ET	11	2.50	2.1		2/0

		1	HLA-A2 Supermotif Pe	ptides		
L2	9	278		E1	8	71
L2	10	278		E1	9	71
L2	11	278		B1	9	178
E6	11	96		E2	8	174
L2	9	356		L2	9	274
L2	10	356		B1	10	250
E7	8	75		B1	8	143
E7	9	75		B2	9	2
E7	10	75		B1	8	21
L2	8	322		E1	10	21
L2	9	322		E2	8	66
L2	9	404		E2	10	66
L2	11	404		L2	8	173
E1	11	570		L2	10	173
E7	8	88		62 81	10	336
E7	11	88		E1	11	336
L2	11	347		E1	11	180
L2	10	396		E1	11	62
L2	11	396				299
		222		B1		100
E1	11	313		L2	11	332
E2	10				8	
L1	8	366		L2		332
L1	11	366				332
E7	8	14			8	192
E7	10	14		L2	9	192
L1	9	208			8	105
L1	11	208			9	105
E1	11	46			11	105
L1	9	195				120
L1	10	195				42
L2	9	42				197
L2	11	42			10	197 197
E6	9	14				
E6	11	14				17
Ll	10	455				17
L1 E2	11	455 141				334 74
E2	11	141				74
L1	10	198				417
E1	9	481				360
E1	11	481				27
E1	10	73				341
L1	8	331				341
L1	9	331				73
L1	10	331				73
L2	10	369				92
L2	11	369				92
E1	8	534				92
LI	10	411				96
L1	11	411				135
L2	10	30				135
L2	11	30				75
E6	9	99				75
L1	9	215				75
L1	10	215				75
L2	11	258				185
L2	8	258 143				185
L2 L2	9	143				39
L2	10	143				141
E2 ·	11	136				141
	11	130				141

				optide		
E1	9	39		L2	10	129
E1	10	39		L2	11	198
E6	8	113		E2	11	171
E6	9	113		E5	8	28
E6	10	113		E5	10	28
L1	9	262		L1	8	326
L1	10	262	•	L1	11	326
L1	11	262		E5	8	24
L1	8	103		E5	9	24
L1	10	103		E5	10	24
E2	8	118		E5	11	24
E2	11	118		L1	8	202
L1	8	381		L1	10	202
L1	9	381		L2	9	117
E7	9	78		L2	10	314
E7	10	78		L1	8	318
E2	8	205		L1	10	318
E2	9	205		L1	11	318
E2	11	205		L2	8	58
E5	8	2		E1	9	243
E5	9	2		E1	11	194
E5	11	2		E1	8	326
L1	11	206		E1	9	326
Ll	8	80		E2	11	156
L1	9	252		E1	9	350
L2	9	442		E1	10	350
L2	11	442		L1	10	101
E1	8	50		L2	10	56
L1	8	369		£7	8	22
L1	9	369		E7	9	22
L1	10	369		E1	8	217
E1	10	454		E2	9	50
L2	9	428		E2	10	50
E5	8	40		L1	9	400
E5	9	40		L1	10	400
E5	10	40		L2	8	292
E5	11	40		L2	10	223
E1	10	494		L1	9	144
L1	8	119		L1	11	144
E1	8	393		E2	8	55
E1	11	393		E2	10	55
E2	10	346		E1	8	273
E2 L2	11	346		L1	8	136
L2	8	398		L1	10	136
E1	9 10	398		E1	10	162
L1	9	446 245		E2	8	162
E1	9			E2	11	162
L2	8	457 239		L1	8	107
L2	9	239		L2	8	300
L2	10	239		E1	8	191
L2	11	276		E1	9	191
E1	11	18		L2	8	215
E2	8	290		L2	10	215
E1	8			L2	8	25
E1	10	252 252		L2	11	25
L1				E1	10	6
Li	8 10	371		L2	9	64
L2	8	371 129		L2	11	64
	9			E1	8	436
L2	9	129		E1	9	436

		HLA-A2 Sup	permotif Peptides		
L2	10	60	L2	11	149
El	11	145	Li	10	361
E7	9	85	E2	8	29
E7	11	85	E1	10	502
L1	10	407	E7	8	5
L2	8	413	E7	9	5
L2	9 '	413	E7	11	5
L2	10	413	E5	8	8
E1	8	467	E5	9	8
E1	9	467	E5	10	8
E1	10	467	E5	11	8
E1	11	467	L2	11	40
E1.	9	147	E1	9	98
L2	8	75	E5	8	22
L2	9	75	E5	10	22
L1	10	222	E5	11	22
E5	8	11	E1	9	474
E5	10	11	L2	8	326
E5	11	11	L2	10	326
E4	8	90	L2	8	287
E4	10	90	E5	9	21
E1	10	316	E5	11	21
L2	9	51	L1	9	272
L2	11	51	L1	11	272
L1	10	111	E5	11	31
L1	10	478	L2	9	113
E2	9	242	L2	10	113
E2	10	242	L2	8	279
L1	8	113	L2	9	279
E1	9	415	. L2	10	279
E1	8	189	E6	10	97
E1	10	189	L2	10	141
E1	11	189	L2	11	141
L1	8	35	E1	10	195
L1	9	35	L2	8	178
L1	10	35	L2	9	178
E2	10	53	E5	10	32
L2	8	177	E5	11	32
L2	9	177	L2	9	44
L2	10	177	E5	9	17
E6	9	119	E5	10	17
E6	11	119	B6	8	120
B1	8	264 264	E6	10	120
E1 E4	11	264 59	L1 E1	9	191 313
E4 E4					
E2	11	59 78	E1 E1	10	265 265
E2	9	310	L2	11	405
E1	10	449	L2	10	405
E2	11	274	L2		429
L2	9	230	L2	11	429
L2	11	230	B1		56
E1	8	230 176	E1	10	56
E1	11	176	E1	11	56
E6	8	25	B1	10	571
L1	8	387	Ll	8	376
E4	10	26	L1	9	376
L2	8	306	L1	11	376
L2	10	306	El	11	341
E1	8	581	L2	11	82
	-				

		1	HLA-A2 Supermotif Pe	ptides		
L2	9	185		L1	11	148
L2	11	185		E2	10	182
L2	8	131		E2	11	182
L2	10	131		E1	11	424
L2	11	131		E2	10	84
L1	10	187		E2	11	84
L1	11	187		E6 -	9	18
E4	9	83		E6	10	18
E4	10	83		E4	8	42
E4	11	83		E1	8	231
		89		E1	9	231
E7	10 11	11		E1	10	231
E7		121		E2	11	115
L2	10			E2	8	
L2	11	121 443		E2	11	165 165
E1 E2	11	287		E1	8	518
E1	8	23		E1	9	518
E1	10	23		L2	8	34
	9	104		L2		34
L2 L2	10	104		E2	11	147
L2	11	104		E2	11	147
		34		E6	9	
E5	9	34		E1	9	116 121
E5		34			11	
E5	11	41		E6 E1		52 283
E5 E5	9	41		E1		283
E5	10	41		E1		283
E5	11	41		E2	10	63
		45			11	63
E2 E1	10 11	553		E2 -	9	61
	8	325		L1	9	19
E2 E2	9	325		L1	10	19
E2	10	325		Ll	11	71
E2	11	325		E1	8	351
L1	9	311		E1		351
	10	123		E1	11	351
L1	11	486		E4	10	13
E1	11	433		E4	9	60
E6	11	73		E4	10	60
E2	10	351		L1		42
E1	8	312		L2		107
E1	9	312		L2	10	107
E2	9	359		E1		255
E2	10	359		E1		255
E1	8	254		E1	11	307
E1	10	254		L1		271
E6	11	128		E1		557
E1	9	357		E1		557
L2	10	22		E1		557
L2	11	22		E5	8	16
E1	9	114		E5		16
E1	8	420		E5		16
Li	8	169		E6		101
L1	10	169		E1		223
E6	8	94		E1		223
E6	9	94		L2	8	179
E7	8	49		B1		491
E2	8	47		E1		491
E2	10	47		E1		491
L1	8	148		E2	9	314
~ *	-				-	

		TIEA-A2 Supermont	cpudes		
L1	11	186	Ll	11	465
L2	8	246	L1	8	209
E5	9	33	L1	10 '	209
E5	10	33	E1	11	132
L1	11	41	E1	8	358
E1	11	521	L2	9	23
E1	9	540	L2	10	23
E2	10	15 208	E4 E6	11	81
E1	11	208 83	E6	9	121 555
E7	8 11	83	E1	10	555
E4	8	8	E1	11	555
E1	9	198	E5	8	49
E1	10	198	E5	9	49
E1	11	198	E5	11	49
E2	8	82	E5	10	70
E7	8	82	E5	11	70
E7	9	82	E1	8	268
E5	11	59	E1	9	268
E5	8	55	E1	10	268
E5	9	55	E1	11	268
E5	11	55	E1	8	115
E5	9	51	L2	8	372
E5	10	51	L2 E6	9	372
E5 E1	11 9	51 298	E5	10 9	38 61
E1	10	298	E5	10	61
E1	11	298	E5	11	61
E5	11	69 -	L2	8	338
E5	10	60	L2	11	338
E5	11	60	E5	8	73
E5	8	72	E5	10	73
E5	9	72	E1	8	514
E5	11	72	B1	9	514
E1	9	276	E7	9	29
E1	9	563	E5	9	47
E1	10	563	E5	10	47
E5	8	56	E5	11	47
E5	10	56	E1	8	277 295
E2 E2	8 10	42 42	L1 E1	8	564
E5	8	42 52	E1	9	564
E5	9	52	E7	8	67
E5	10	52	E7	10	67
E5	11	52	L1	8	95
E2	8	94	L1	10	95
E2	11	94	L1	10	233
E5	8	65	E4	8	1
E5	9	65	E4	9	1
E5	10	65	E4	11	1
Ll	10	367	L2	8	87
L1	11	367	L2	11	87
E6	11	27	L1	11	383
L1	11	309	E1	8	306
E1	9	511	B1	10	398
E1	10	511	E1 E2	11 8	398 75
E1 E7	11 9	511 15	E2	9	75
L2	8	399	E2	11	75
L2	11	399	E2	9	56
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			HLA-AZ Supermotii	Peptide	s	
L1	8	338		L1	9	142
L1	9	338		Ll	11	142
E2	10	151		E7	9	64
E1	10	47		E7	11	64
E1	11	47		E2	8	348
L1	8	196		E2	9	348
L1	9	196		L2	10	237
E1	10	19		L2	11	237
L1	8	154		L2	8	124
E1	11	274		L2	9	124
E1 L2	10 8	361 115		L2	9	285
L2	11	115		L2 L2	10 8	285 139
E2	9	71		L2	9	139
E2	8	249		E5	9	78
E2	11	249		E5	10	78
E6	8	36		E2	8	216
E1	11	607		E2	8	196
L2	8	270		E2	11	196
L2	10	270		L1	8	482
L2	11	270		L1	9	482
E1	10	389		L2	9	325
E6	8	5		L2	11	325
E6	9	5		L1	8	217
E1	9	329		L2	9	189
El	9	600		L2	11	189
E1	8	270		E1	8	94
E1	9	270		E1	9	94
El	10	270		E1	11	442
E1	11	270		E6	8	110
E1	8	451		E6	11	110
Ll	11	31		E4	10	34
E1 E1	8 9	300 300		L1 L1	8	183
L2	8	366		L1	9	183
L2	10	366		L2	9	451 451
E1	9	55		L1	8	458
E1	11	55		L2	10	73
L1	10	445		L2	11	73
E1	8	539		E7	8	47
E1	10	539		E7	و	47
L1	9	438		E7	10	47
E6	9	21		E1	8	562
El	11	397		El	10	562
L1	8	337		E1	11	562
Ll	9	337		E5	8	64
L1	10	337		E5	9	64
L1	В	323		E5	10	64
L1	10	323		E5	11	64
Ll	11	323		E1	8	258
E1	10	304		E1	11	258
E6	9	75		L2	9	389
L2	8	38		L2	11	389
E1	11	96		L2	9	337
E2	9	127		E1	8	513
E2 L1	11	127 289		E1 E1	9 10	513
L2	9	385		L2	8	513 86
L2	10	377		L2	9	86
L2	11	377		L2	9	411
		J.,			-	411

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L2	10	411		L1	9	419
L2	11	411		E1	9	399
E1	10	545		E1	10	399
E1	11	545		E1	11	399
L2	11	168		E1	9	64
L2	11	243		E1	11	64
L2	8	423		E5	9	7
E2	10	354		E5	10	7
L2	8	183		E5	11	7
L2	11	183		L2	8	43
E4	8	67		L2	10	43
E4	10	67		E6	10	28
E4	11	67		E6	11	28
E1	9	182		E1	10	31
E1	11	182		E6	8	15
L2	9	359		E6	10	15
L2	10	207		L1	8	151
L2	11	207		L1	11	151
	9	90		L1	9	372
L1					11	
L1	11	90		L1		372
L2	9	96		L1	9	301
L2	10	96		L1	9	324
E2	9	258		L1	10	324
E2	10	258		E2	11	14
E2	11	258		E7	9	81
L2	8	171		E7	10	81
L2	9	171		E7	10	28
L2	10	171		L2	8	16
L2	11	426		L2	9	16
L2	8	158		L2	11	16
L2	9	158		E4	8	4
E7	10	20		E4	10	4
E7	11	20		E4	11	4
E5	8	19		L1	11	232
E5	11	19		L1	11	250
L1	8	266		E2	9	48
L2	11	212		E2	11	48
L1	10	175		E2	8	76
E5	8	5		E2	10	76
E5	9	5		E1	9	305
E5	11	5		E2	9	344
L1	8	16		E7	8	80
L1	10	16		E7	10	80
E2	10	222		E7	11	80
E2	11	222		E2	8	244
L2	10	418		E2	10	244
L2	8	363		E2	11	244
L2	10	363		L2	8	19
L2	11	363		L1	9	210
L2	8	91		L1	11	210
L2	8	252		E2	8	233
L2	10	252		E2	11	233
L2	8	328		L1	10	149
L2	11	328		E2	10	218
E1	9	636		B1	8	344
E1	10	636		E1	9	344
E1	11	636		L2	8	231
L1	8	177		L2	10	231
L1	10	177		L2	8	233
L1	11	177		L2		233

		HLA-AZ Su	permour re	pudes		
E2	8	57		L2	8	111
E2	11	57		L2	11	111
E1 .	8	391		L2	11	77
E1	10	391		E6	8	3
L1	9	259		E6	9	3
L2	8	227		E6	10	3
L2	9	227		E6	11	3
L1	10	53		L2	10	181
L2	8	5		L2	8	13
L2	10	5		L2	11	13
L2	11	5		E6	8	9
L2	10	11		E6	11	9
L2	8	298		E1	8	547
L2	10	298		B1	9	547
L2	8	316		E1	10	547
L2	11	316		E1	11	547
L2	9	449		L2	9	153
L2	10	449		L2	9	267
L2	11	449		L2	11	267
E2	9	7		E5	8	30
E2	10	7		L1	8	50
E1	8	109		L1	9	50
L1	11	241		L1	10	50
E1	8	125		E2	9	207
E1	9	125		E2	11	207
L2	8	281		L2	8	3 92
L2	9	245		L1	10	474
E2	9	303		L1	11	474
E1	8	616		E1	9	247
E7	9	66		E1	10	247
E7 L1	11 9	66 94		L1 L1	9	375 375
L1	11	94		L1	10	375
E1	9	69		L2	8	81
E1	10	69		L2	10	103
E1	11	69		L2	11	103
E1	10	117		L1	11	285
E2	8	343		L1	10	86
E2	10	343		L1	11	86
E1	9	343		L2	9	49
E1	10	343		L2	11	49
E1	9	324		L2	8	106
E1	10	324		L2	9	106
E1	11	324		L2	11	106
L1	8	476		E1	9	490
L1	9	476		E1	10	490
L2	9	68		E1	11	490
L2	11	68		E2	9	81
E1	9	293		E4	8	80
L1	8	29		L1	9	294
L1	8	279		E1	9	260
L1	10	279		E1	10	260
L2	8	221		E2	10	88
L2	9	221		E6	10	23
Ll	11	140		L2	10	304
L1	9	488		E2	8	150
Ll	11	488		E2	11	150
L1	8	379		E1	10	635
L1	10	379		E1	11	635
L1	11	379		L1	10	418

		TICA-A2 Superino	an repnaes		
E1	8	354	E1	10	97
E1	9	354	E6	8	12
E7	10	45	E6	11	12
E7	11	45	E2	9	355
E2	8	23	E2	11	355
E2	9	23	L2	10	184
E4	8	86	L2	8	130
E4	9	86	L2	9	130
E4	10	86	L2	11	130
E5	8	14 14	E4 E4	10	82 82
E5	9 10	14	E1	11 8	294
E5 L1	8	335	Li	8 .	339
L1	10	335	L1	11	339
L1	11	335	E7	8	86
L2	8	241	E7	10	86
L2	8	210	E4	9	68
E2	8	220	E4	10	68
E2	9	226	E4	11	68
E6	10	7	Ll	9	408
E2	8	201	L1	11	270
E2	9	211	E1	8	556
E2	11	211	E1	9	556
E1	8	289	E1	10	556
E1	9	289	E1	11	556
E1	10	289	E5	8	15
E1	11	289 190	E5 E5	9 11	15 15
E2 E2	8 10	190	E7	9	7
E1	11	331	E5	8	50
L1	11	7	E5	10	50
E2	10	317	E5	11	50
ь1	8	281	E1	10	297
L1	10	281	E1	11	297
L1	8	189	E5	9	71
L1	9	189	E5	10	71
Ll	11	189	E2	9	93
L1	10	392	L1	8	377
E2	10	40	Ll	10	377
E5	8	45	E1	9	408
E5	9	45	E1 E2	11 8	408 128
E5	11 9	45 260	E2	10	128
L2 E1	8	185	E2	8	227
E1	9	185	E2	10	203
E1	11	185	E2	11	203
E2	9	198	L2	8	360
E2	11	198	L2	11	360
L2	9	164	E1	11	30
L2	11	164	L1	9	150
L2	8	145	L2	9	15
L2	10	145	L2	10	15
L1	9	343	E1	8	526
E6	8	40	E1	10	526
El	8	192	L2	9	166
L2	8	420	L2	8	380
L2	10	420	L2 L2	9 11	380
L2	11	420 409	L2 L1	9	92
L2 L2	9	216	L1	11	92
112	-	210	-		

		ni.A-A2 Super	mom reputes		
E1	8	148	L2	10	378
E6	9	39	L2	11	378
E1	8	232	E4	8	92
E1	9	232	L1	9	328
L1	9	223	L1	11	328
L1	11	223	L1	10	8
E6	8	142	E1	9	317
E6	9	11	E1.	11	317
L2	8	137	L2	10	339
L2	10	137	E1	10	239
L2	11	137	E1	8	519
E5	8	62	L2	8	97
E5	9	62 .	L2	9	97
E5	10	62	L2	11	97
E5	11	62	B1	8	291
E2	11	202	E1	9	291
L1	8	91	B1	11	291
L1	10	91	L1	8	21
L1	8	332	L1	10	21
L1 L1	9 11	332 332	E2 E2	8 11	192 192
	9	151	B2 B1	9	
L2 L2	11	151	E1	10	333
E4	8	77	E5	10	20
E4	9	77	L2	9	31
E4	11	77	L2	10	31
E4	8	84	L2	11	31
E4	9	84	Li	8	190
E4	10	84	Li	10	190
E4	11	84	L2	10	199
E5	9	12	E7	10	12
E5	10	12	L1	9	393
E5	11	12	E6	10	53
L2	9	136	E6	11	53
L2	11	136	E4	8	7
L2	10	150	E4	9	7
E4	8	76	E1	10	275
E4	9	76	E2	9	41
E4	10	76	E2	11	41
E6	11	87	L1	9	73
L2	8	386	L1	10	73
E4	9	91	E5	8	48
E2	8	212	E5	9	48
E2	10	212	E5	10	48
E1	8	290	E5	8	46
E1	9	290	E5	10	46
E1	10	290	E5	11	46
E6	10	88	Ll	8	382
E6	11	88	L1	9	176
E4	9	73	L1	11	176
E4	10	73	E7	8	69
E4	11	73	E7	11	69
E2	10	116	E1	9	79
E2	9	191	E2 E2	8 10	139 139
E1	8 10	345	E2 E1	9	444
E1		332	E2	10	
E1	11	332	L2	8	288
L2	11	387 78	L2 L2	11	261
E1 L2	10 9	378	E1	9	24
<u>ы</u> г .	,	3/0	E1	,	44

Table VIIIA HPV 6A HLA-A2 Supermotif Peptides

		III.A-Az Superinour i	epitics		
E5	8	6	E1	11	464
E5	10	6	E5	8	58
E5	11	6	E1	8	510
E7	8	79	B1	10	510
E7	9	79	E1	11	510
E7	11	79	E1	8	267
E2	8	243	E1	9	267
E2	9	243	E1	10	267
E2	11	243	E1	11	267
E2	9	232	E2	9	134
L2	9	232	E2	10	92
L2	11	232	E6	8	141
E1	9	362 .	E6	9	141
L2	9	419	E4	10	72
L2	11	419	E4	11	72
E7	11	55	E2	8	145
L2	9	234	E2	10	145
E7	8	6	E1	8	237
E7 -	10	6	E6	8	61
L2	9	364	L2	9	349
L2	10	364	E6	8	82
L2	8	165	E1	8	262
L2	10	165	E1	10	262
L2	8	379	E1	11	380
L2	9	379	E6	9	85
L2	10	379	E6	8	46
L1	10	27	E6	9	46
L2	9	146	E5	9	81
E1	8	565	L1	9	385
L1	8	344 327	L1 E4	10 11	385 12
L1 E1	10 11	238	E6	10	105
L1	8	20	E1	10	86
L1	9	20	L2	9	435
L1	11	20	E1	8	579
E5	8	25	E1	10	579
E5	9	25	E5	8	54
E5	10	25	E5	9	54
E5	11	25	E5	10	54
L1	8	329	L2	8	371
L1	10	329	L2	9	371
L1	11	329	L2	10	371
E2	8	349	E1	9	532
L1	10	72	E1	10	532
L1	11	72	L1	10	358
E2	10	58	E1	11	536
E5	8	3	L2	9	18
E5	10	3	L1	8	65
E5	11	3	L1	9	65
E7	9	68	L1	10	65
E2	8	132	E2	8	159
E2	11	132	E2	11	159
L1	8	97	L1	10	350
E2	11	321	E2	8	214
E1	9	426	E2	10	214
E5	9	36	E5	8	43
E5	11	36	E5	9	43
E1	8	340	E5	10	43
E1	8	530	E5 E1	11 8	43 402
E1	11	530	E-T	0	402

E4	8	6
E4	9	6
E4	10	6
E2	8	168
Ll	9	287
L1	10	287
L2	8	71
Ll	8	10
L1	11	10
E2	9	138
E2	11	138
L1	8	415
E1	8	91
E1	9	91
E1	11	91
L1	11	26
E2	9	131

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2	3	4	E4	8	71
L1	9	234	E4	9	71
L2	10	329	L2	10	14
L2	11	329	L2	11	14
ESA	9	9	E1	9	525
E5A	10	9	E1	11	525
E1	8	318	E6	10	10
E1	10	318	E6	8	86
L1	8	489	E1	11	77
L1	10	489	E1	10	101
L2	9	340	L1	9	43
L2	11	340	E5B	8	36
E4	8	12	E2	10	231
E4	10	12	L1	8	483 .
E2	8	3	E1	8	601
L2	8	286	E6	11	64
L2	9	286	E5B	8	20
E2	8	72	E5B	10	20
E2	11	72	E5B	11	20
L2	10	112	E2	9	124
L2	11	112	E2	10	124
E1	11	112	Li	9	341
L2	8	140	L1	11	341
L2	11	140	E1	11	406
L1	8	420	E1	10	473
E1	8	475	E1	11	234
E1	9	22	E6	8	67
E1	11	22	E6	9	137
E2	10	250	E2	9	296
E1	8	65	E2	8	35
E1	10	65	E2	9	35
E4	9	24	E1	8	488
E4	11	24	E1	11	488
L2	8	228	L1	9	153
L2	11	228	E2	8	11
Li	11	81	E5B	8	5
L2	9	421	E5B	10	5
L2	10	421	E7	9	71
E1	10	554	E1	9	14
E1	11	554	E1	10	14
E6	11	37	E1	11	14
E5A	8	79	Li	8	171
E5A	9	79	E6	8	131
E5A	10	79	E2	8	252
E5A	11	79	E6	8	31
E1	9	319	E6	9	31
L1	9	203	E6	10	31
L2	9	327	E6	11	31
E1	10	63	E1	9	640
L2	8	341	E1	10	640
L2	10	341	E4	10	74
L1	8	312	E4	11	74
L1	10	300	E2	8	9
E4	9	13	E2	10	9
E4	11	13	E2	8	153
E1	10	381	E1	10	516
E2	11	217	E1	11	516
L1	9	22	E1	8	524
L1	11	22	E1	10	524
E1	11	296	E2	11	230
E1	10	407	L1	9	24
101	70	407		-	

E1	8	369		L2	10	143
E1	9	369		E2	8	348
E1	8	170		E2	9	348
E1	10	170		E2	9	136
L2	9	278		E2	11	136
L2	10	278		E1	8	71
L2	11	278		E1	9	71
E6	11	96		E1	9	178
E7	8	75		E2	8	174
E7	9	75		L2	9	274
E7	10	75		E1	10	250
L2	8	322		E1	8	143
L2	9	322		E2	9	2
L2	9	403		E1	8	21
L2	11	403		E1	10	21
E1	11	570		E2	8	66
L2	11	347		E2	10	66
L2	10	395		L2	8	173
L2	11	395		L2	10	173
E1	11	222		E1	10	336
E2	10	313		E1	11	336
				E1		
L1	8 11	366 366		E1	11 11	180 62
L1						
E7	8	14		L1	11	299
E7	10	14		E1	11	100
Ll	9	208		L2	8	332
L1	11	208		L2	9	332
E1	11	46		L2	10	332
Ll	9	195		L2	8	192
L1	10	195		L2	9	192
L2	9	42		E1	8	105
L2	11	42		E1	9	105
E6	9	14		E1	11	105
E6	11	14		L2	11	120
L1	10	455		E6	10	42
L1	11	455		E1	8	197
L1	10	198		E1	10	197
E1	9	481		E1	11	197
E1	11	481		E2	8	17
E1	10	73		E2	10	17
L1	8	331		L2	8	334
L1	9	331		E2	9	74
L1	10	331		E2	10	74
E5B	8	11		E1	11	417
E5B	9	11		E1	11	360
E5B	11	11		E7	11	27
E2	9	143		E2	9	341
E2	10	143		E2	10	341
L2	10	369		E7	10	73
L2	11	369		E7	11	73
E1	8	534		E6	9	92
L1	10	411		E6	10	92
L1	11	411		E6	11	92
L2	10	30		E2	9	96
L2	11	30		L2	8	135
E6	8	99		L2	10	135
L1	9	215		B4	8	85
L1	10	215		E4	9	85
L2	11	258		E4	10	85
L2	8	143		E4	11	85
L2	9	143		E2	8	185

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E2	9	185	B1	9	457
E7	10	39	L2	11	276
E1	8	141	E1	11	18
E1	10	141	E2	8	290
E1	9	39	E1	8	252
E1	10	39	E1	10	252
E6	8	113	L1	8	371
E6	9	113	L1	10	371
E6	10	113	L2	8	129
L1	9	262	L2	9	129
L1	10	262	L2	10	129
LI	11	262	E2	11	171
L1	8	103	E5A	8	28
LI	10	103	E5A	10	28
E2	8	118	L1	8	326
E2	11	118	L1	11	326
L1	8	381	E5A	8	24
L1	9	381	E5A	9	24
E7	9	78	B5A	10	24
E7	10	78	E5 A	11	24
E2	8	205	L2	11	198
E2	9	205	L1	8	202
E2	11	205	L1	10	202
E5A	8	2	L2	9	117
E5A	9	2	B4	9	2
E5A	11	2	E4	10	2
L1	11	206	E4	11	2
L1	8	80	L2	10	314
L1	9	252	L1	8	318
L2	9	442	L1	10	318
	11	442	Ll	11	318
L2	8	50	L2	8	58
E1		369	E1	9	243
L1 L1	9	369	E1	11	194
		369	L2	9	356
L1	10	16	L2	10	356
E5A			E1		
E5A	10	16	E1	8	326
E5A	11	16		9	326
E1	10	454	E2	11	156
L2	9	428	E1	9	350
E5B	8	22	E1	10	350
E5B	9	22	L1	10	101
E5B	10	22	L2	10	56
E5B	11	22	E7	8	22
E5A	8	40	E7	9	22
E5A	9	40	E5B	9	28
E5A	10	40	E5B	10	28
E5A	11	40	E1	8	217
E2	10	346	E2	9	50
E2	11	346	E2	10	50
E1	10	494	L1	9	400
L1	8	119	L1	10	400
E1	8	393	L2	8	292
E1	11	393	E5B	9	15
L2	8	397	E5B	10	15
L2	9	397	L2	10	223
E1	10	446	L1	9	144
L1	9	245	L1	11	144
L2	8	239	E5B	8	25
L2	9	239	E2	8	55
L2	10	239	E2	10	55

PCT/US00/33549

			TIDIT TIE Dapermont	*pildes		
E1	8	273		E2	8	78
L1	8	136		E2	9	310
L1	10	136		E1	10	449
El	10	162		E1	11	449
E2	8	162		E2	11	274
E2	11	162		L2	9	230
L1	8	107		L2	11	230
L2	8	300		E1	8	176
E1	8	191		E1	11	176
E1	9	191		E6	8	25
L2	8	215		L1	8	387
L2	10	215		E4	10	36
L2	8	25		L2	8	306
L2	11 .	25		L2	10	306
E1	10	6		E1	8	581
L2	9	64 64		L2 L1	11	149
L2	11	436		E2	10	361 29
E1	9	436		E1	10	502
E1 L2	10	60		E7	8	502
E1	11	145		E7	9	5
L1	10	407		E7	11	5
E7	9	85		E5A	8	8
E7	11	85 .		E5A	10	8
L2	8	412		E5A	11	8
L2	10	412		L2	11	40
L2	11	412		E1	9	98
E1	8	467		E5A	8	22
E1	9	467		E5A	10	22
E1	10	467		ESA	11	22
E1	11	467		E1	9	474
E1	9	147		L2	8	326
L1	10	222		L2	10	326
E5A	8	11		L2	8	287
E5A	10	11		E5A	9	21
E5A	11	11-		E5A	11	21
E4	8	100		L1	9	272
E4	10	100		L1 E5A	11	272
E1	10 9	316 51		L2	11 9	31 113
L2 L2	11	51		L2	10	113
L1	10	111		L2	8	279
L1	10	478		L2	9	279
E2	9	242		L2	10	279
E2	10	242		E6	10	97
L1	8	113		L2	10	141
E1	9	415		L2	11	141
E1	8	189		E1	10	195
E1	10	189		L2	8	178
E1	11	189		L2	9	178
L1	8	35		E5A	10	32
L1	9	35		E5A	11	32
L1	10	35		L2	9	44
E2	10	53		E5A	9	17
L2	8	177		E5A	10	17
L2	9	177		E6	8	120
L2	10	177		E6	10	120
E6	9	119		L1	9	191
E6	11	119		E1	8	313
E1	8	264		E1 E1	10 11	265 265
E1	11	264		E1	11	∠65

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L2	8	404	E1	9	357
L2	10	404	L2	10	22
L2	8	429	L2	11	22
L2	11	429	E1	9	114
E1	8	56	E1	8	420
E1	10	56	L1	8	169
E1	11	56	L1	10	169
E1	10	571	E6	8 .	94
LI	8	376	E6	9	94
L1	9	376	E7	8	49
1.1	11	376	E2	8	47
El	11	341	E2	10	47
L2	11	82	L1	8	148
L2 L2	8 10	131 131	L1 E1	11	148 424
	11	131	E6	9	18
L2					
L1	10	187	E6	10	18
L1	11	187	E4	8	52
E4	9	93	E1	8	231
E4	10	93	E1	9	231
E4	11	93	E1	10	231
E7	10	89	E2	11	115
E5B	8	23	E2	8	165
E5B	9	23	E2	11	165
E5B	10	23	E1	8	518
E7	11	11	E1	9	518
L2	10	121	L2	8	34
L2	11	121	L2	11	34
E1	10	443	E2	8	147
E2	11	287	E2		147
E1	8	23	E6	9	116
E1	10	23	E1	9	121
L2	9	104	E6	11	52
L2	10	104	E1	8	283
L2	11	104	E1	9	283
E5A	8	34	E1	10	283
E5A	9	34	E2	10	63
E5A	11	34	E2	11	63
E5A	8	41	L1	9	61
E5A	9	41	L1	9	19
E5A	10	41	L1	10	19
E5A	11	41	L1	11	71
E2	10	45	E1	8	351
E1	11	553	E1	9	351
E2	8	325	E1	11	351
E2	9	325	E4	10	23
E2	10	325	E4	9	70
E2	11	325	E4	10	70
L1	9	311	L1	10	42
E6	10	123	L2	8	107
L1	11	486	L2	10	107
E1	11	433	E1	9	255
E6	11	73	E1	11	255
E2	10	351	E1	11	307
E1	8	312	L1	10	271
E1	9	312	El	8	557
E2	9	359	E1	9	557
E2	10	359	E1	10	557
E1	8	254	E6	11	101
E1	10	254	E1	10	223
E6	11	128	B1	11	223
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L2	В	179	L1	11	367
E1	В	491	E6	11	27
E1	9	491	L1	11	309
E1	10	491	E1	9	511
E2	9	314	E1	10	511
L1	11	186	B1	11	511
L2	8	246	E7	9	15
E5A	9	33	L2	8	398
E5A	10	33	L2	11	398
L1	11	41	L1	11	465
E5B	9	18	L1	8	209
E5B	10	18	L1	10	209
E1	11	521	L2	9	74
E1	9	540	L2	10	74
E2	10	15	E5B	10	3
E1	11	208	B1	11	132
E7	8	83	E1	8	358
E7	11	83	L2	9	23
E4	8	18	L2	10	23
E1	9	198	E4	11	91
E1	10	198	E6	9	121
E1	11	198	B1	8	458
E7	8	82	E1	11	458
E7	9	82	E1	9	555
E5B	8	29	E1	10	555
E5B	9	29	B1	11	555
E5A	11	59	E5A	8	49
E5A	В	55	E5A	9	49
E5A	9	55	E5A	11	49
E5A	11	55 -	B5A	10	70
E5A	9	51	E5A	11	70
E5A	10	51	E1	8	268
E5A	11	51	E1	9	268
E5B	8	30	B1	10	268
E1	9	298	E1	11	268
E1	10	298	E1	8	115
B1	11	298	L2	8	372
E2	8	82	L2	9	372
E5A	11	69	E6	10	38
E5A	10	60	E5A	9	61
E5A	11	60	E5A	10	61
		72	E5A	11	61
E5A	В		L2		
E5A	9	72		8	338
E5A	11	72	L2	11	338
E1	9	276	E5A	8	73
E1	9	563	E5A	10	73
E1	10	563	E1	8	514
E5A	8	56	E1	9	514
E5A	10	56	E7	9	29
E2	В	42	E1	8	277
E2	10	42	E5A	9	47
E5A	8	52	E5A	10	47
E5A	9	52	E5A	11	47
E5A	10	52	L1	8	295
E5A	11	52	E2	10	222
E2	В	94	E2	11	222
E2	11	94	E1	8	564
E5A	8	65	E1	9	564
E5A	9	65	E7	8	67
E5A	10	65	E7	10	67
L1	10	367	Ll	8	95

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L1	10	95	E1	8	539
L1	10	233	E1	10	539
E4	8	11	L1	9	438
E4	9	11	E6 .	9	21
E4	11	11	E1	11	397
L2	8	87	L1	8	337
L2	11	87	L1	9	337
L1	11	383	L1	10	337
E5B	11	26	Ll	8	323
E1	8	306	L1	10	323
E5B	11	2	L1	11	323
E1	10	398	E1	10	304
E1	11	398	E6	9	75
E2	8	75	L2	8	38
E2	9	75	E1	11	96
E2	11	75	E2	9	127
E2	9	56	E2	11	127
L1	8	338	E1	11	607
L1	9	338	L1	8	289
E2	10	151	L2	8	385
E1	10	47	L2	10	377
E1	11	47	L2	11	377
L1	8	196	L1 L1	9 11	142 142
L1	9	196 19	E7	9	64
E1 L1	10	154	E7	11	64
E1	8 11	274	L2	10	237
E1	10	361	L2	11	237
L2	8	115	L2	8	124
L2	11	115	L2	9	124
E2	9	71	L2	9	285
E2	8	249	L2	10	285
E2	11	249	L2	8	139
E6	8	36	L2	9	139
L2	8	270	L2	8	420
L2	10	270	L2	10	420
L2	11	270	L2	11	420
E1	10	389	E5A	9	78
E6	8	5	E5A	10	78
E6	9	5	E5A	11	78
B1	9	329	E2	8	216
E1	9	600	E2	8	196
E1	8	270	E2	11	196
E1	9	270	L1	8	482
E1	10	270	L1	9	482
E1	11	270	L2	9	325
E1	8	451	L2	11	325
E1	9	451	Ll	8	217 189
L1	11	31	L2 L2	9 11	189
E4	9	5 5	E1	8	94
E4		5	E1	9	94
E4	11	300	E1	11	442
E1 E1	B 9	300	E4	10	69
L2	8	366	E4	11	69
L2	10	366	E6	8	110
E1	9	55	E6	11	110
E1	11	55	E4	10	44
E7	8	88	L1	8	183
E7	11	88	Li	9	183
L1	10	445	L2	8	451
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L2	9	451	L1	10	175
Ll	8	458	E5A	8	5
E7	8	47	E5A	9	5
E7	9	47	E5A	11	5
E7	10	47	Ll	8	16
E1	8	562	L1	10	16
E1	10	562	L2	8	363
E1 .	11	562 64	L2 L2	10	363
E5A E5A	9	64	L2	8	363 417
E5A	10	64	L2	11	417
E5A	11	64	L2	8	91
L2	10	73	L2	8	252
L2	11	73	L2	10	252
L2	8	389	L2	8	328
L2	10	389	L2	11	328
E1	8	258	E1	9	636
E1	11	258	E1	10	636
L2	9	337	E1	11	636
E1	8	513	Ll	8	177
E1	9 10	513	L1 L1	10 11	177 177
E1 L2	8	513 86	L1	9	419
L2	9	86	E1	9	399
L2	9	410	E1	10	399
L2	10	410	El	11	399
E1	10	545	El	9	64
E1	11	545	E1	11	64
L2	11	168	E5A	9	7
L2 '	11	243	E5A	11	7
L2	8	423	L2	8	43
E2	10	354	L2	10	43
E4	8	77	E6	10	28
E4	10	77	E6	11	28
E4 E1	11 9	77 182	E1 E6	10 8	31 15
E1	11	182	E6	10	15
L2	9	359	Li	8	151
L2	10	207	Li	11	151
L2	11	207	Li	9	372
L1	9	90	Ll	11	372
L1	11	90	L1	9	301
L2	8	183	L1	9	324
L2	11	183	L1	10	324
L2	9	96	E2	11	14
L2	10	96	E7	9	81
E2 E2	9 10	258 258	E7 E7	10	81 28
E2	11	258	L2	8	16
L2	8	171	L2	9	16
L2	9	171	L2	11	16
L2	10	171	E4	8	14
L2	11	426	E4	10	14
L2	8	158	E4	11	14
L2	9	158	Ll	11	232
E7	10	20	Ll	11	250
E7	11	20	E2	9	48
E5A	8	19	E2	11	48
E5A	11	19	E2	8	76
L1 L2	8 11	266 212	E2 E1	10 9	76 305
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Table VIIIB HPV 6B HLA-A2 Supermotif Peptides

			TIEA-A2 Supermot	ii i cpiace		
E2	9	344		E2	10	343
E7	8	80		E1	9	343
E7	10	80		E1	10	343
E7	11	80		E2	10	84
E2	8	244		E2	11	84
E2	10	244		E1	9	324
E2	11	244		El	10	324
L2	8	19		E1	11	324
L1	9	210		L1	8	476
Ll	11	210		L1	9	476
E2	8	233		L2	9	68
E2	11	233		L2	11	68
L1	10	149		E1	9	293
L2	8	75		L1	8	29
L2	9	75		L1	8	279
E2	10	218		L1	10	279
E1	8	344		L2 L2	8 9	221
E1	9	344 231		L1	11	140
L2 L2	10	231		L1	9	488
L2	8	233		Li	11	488
L2	10	233		Ll	8	379
E2	8	57		L1	10	379
E2	11	57		L1	11	379
E1	8	391		L2	8	111
E1	10	391		L2	11	111
L1	9	259		L2	11	77
L2	8	227		E6	8	3
L2	9	227		E6	9	3
L1	10	53		E6	10	3
L2	8	5		E6 L2	11	3
L2 L2	10 11	5 5		L2	10 8	181 13
L2	10	11		L2	11	13
E5B	9	35		E6	8	9
L2	8	298		E6	11	وَ
L2	10	298		E1	8	547
L2	8	316		E1	9	547
L2	11	316		E1	10	547
L2	9	449		El	11	547
L2	10	449		L2	9	153
L2	11	449		L2	9	267
E2	9	7		L2	11	267
E2	10 8	7 109		E5A L1	8	30 50
E1 L1	11	241		L1	9	50
E1	8	125		L1	10	50
E1	9	125		E2	9	207
L2	8	281		E2	11	207
L2	9	245		L1	10	474
E2	9	303		L1	11	474
E1	8	616		El	9	247
E7	9	66		El	10	247
E7	11	66		Ll	8	375
L1	9	94		L1	9 10	375 375
L1	11	94 69		Ll L2	8	81
El El	9 10	69		L2	10	103
E1	11	69		L2	11	103
E1	10	117		Ll	11	285
E2	8	343		L1.	10	86

WO 01/41799

PCT/US00/33549

L1	11	86	E2	10	40
L2	9	49	E5A	8	45
L2	11	49	E5A	9	45
L2	8	106	E5A	11	45
L2	9	106	L2	9	260
L2	11	106	E1	8	185
E1	9	490	El	9	185
E1	10	490	E1	11	185
E1	11	490	E2	9	198
E2	9	81	E2	11	198
L2	8	391	L2	9	164
E4	8	90	L2	11	164
L1	9	294	L2	8	145
E1	9	260	L2	10	145
E1	10	260	L1	9	343
E2	10	88	E6	8	40
E6	10	23	E1	8	192
L2	10	304	L2	11	408
E2	8	150	L2	9	216
E2	11	150	E1	10	97
E1	10	635	E6	8	12
E1	11	635	E6	11	12
L1	10	418	E2	9	355
E1	8	354	E2	11	355
E1	9	354	L2	8	130
E7	10	45	L2	9	130
E7	11	45	L2	11	130
E2	8	23	E4	10	92
E2	9	23	E4	11	92
E5A	8	14	E1	8	294
E5A	9	14	L1	8	339
E5A	10	14	L1	11	339
E4	8	96	E4	9	78
E4	9	96	E4	10	78
E4	10	96	E4	11	78
E2	9	226	L1	9	408
E2	8	220	Ll	11	270
L1	8	335	E1	8	556
L1	10	335	E1	9	556
L1	11	335	E1	10	556
L2	8	241	E1	11	556
L2	8	210	E7	9	7
E6	10	7	E5A	8	50
E2	8	201	E5A	10	50
E2	9	211	E5A	11	50
E2	11	211	E1	10	297
E1	8	289	E1	11	297
E1	9	289	E5A	9	71
E1	10	289	E5A	10	71
E1	11	289	E7	8	86
E2	8	190	E7	10	86
E2	10	190	E2	9	93
El	11	331	Ll	8	377
L1	11	7	L1	10	377
E2	10	317	E2	11	221
L1	8	281	E1	9	408
L1	10	281	E1	11	408
L1	8	189	E2	8	128
L1	9	189	E2	10	128
L1	11	189	E5B	11	59
L1	10	392	E2	10	203

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E2	11	203		E1	8	290
L2	8	360		E1	9	290
L2	11	360		E1	10	290
E1	11	30		E6	10	88
L1	9	150		E6	11	88
L2	9	15		E4	9	83
L2	10	15		E4	10	83
E1	8	526		E4	11	83
E1	10	526		E2	10	116
L2	9	166		E2	9	191
L2	8	380		E1	8	345
L2	9	380		E1	10	332
L2	11	380		E1	11	332
L1	9	92		L2	10	387
L1	11	92		E1	10	78
E1	8	148		L2	9	378
E6	9	39		L2	10	378
L1	9	223		L2	11	378
Ll	11	223		L2	10	184
E1	8	232		E4	8	102
E1	9	232		L1	9	328
E6	8	142		L1	11	328
E5B	9	63		Ll	10	8
E6	9	11		E1	9	317
L2	8	137		E1	11	317
L2	10	137		L2	10	339
L2	11	137		E1	10	239
E5A	8	62		E1	8	519
E5A	9	62		L2	8	97
E5A	10	62		L2	9	97
E5A	11	62		L2	11	97
E2	11	202		E1	8	291
L1	8	91		E1	9	291
L1	10	91		E1	11	291
L1	8	332		L1	8	21
L1	9	332		L1	10	21
L1	11	332		E2	8	192
L2	9	151		E2	11	192
L2	11	151		E1	9	333
E4	8	87		E1	10	333
E4	9	87		E5A	10	20
E4	11	87		L2	9	31
E5A	9	12		L2	10	31
E5A	10	12		L2	11	31
E5A	11	12		L1	8	190
E4	8	94		L1	10	190
E4	9	94		E7	10	12
E4	10	94		L1	9	393
E4	11	94		E6	10	53
L2	9	136		E6	11	53
L2	11.	136		E4	8	17
L2	10	150		E4	9	17
E4	8	86		E1	10	275
E4	9	86		E2	9	41
E4	10	86		E2	11	41
Еб	11	87		L1	9	73
L2	11	386		L1	10	73
E4	9	101		E5A	8	48
E2	8	212		E5A	9	48
E2	10	212		E5A	10	48
E2	11	212		E5A	8	46

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E5A	10	46	L1	10	72
E5A	11	46	L1	11	72
E4	8	10	E2	10	58
E4	9	10	E5A	8	3
E4	10	10	E5A	10	3
Li	8	382	E5A	11	3
L1	9	176	E7	9	68
L1	11	176	L2	10	199
E7	8	69	E2	8	132
E7	11	69	L1	8	97
E1	9	79	E2	11	321
E2	8	139	E1	9	426
E2	10	139	E5A	9	36
E1	9	444	E5A	11	36
E2	10	288	E1	8	340
L2	8	261	E1	8	530
L2	11	261	E1	11	530
E1	9	24	E5B	9	13
E5A	8	6	E5B	11	13
E5A	10	6	E1	11	464
E7	8	79	E5B	8	17
E7	9	79	E5B	10	17
E7	11	79	E5B	11	17
E2	8	243	E5A	8	58
E2	9	243	E1	8	510
E2	11	243	E1	10	510
E2	9	232	E1	11	510
L2	9	232	E1	8	267
L2	11	232	E1	9	267
E1	9	362	E1	10	267
E7	11	55	E1	11	267
L2	9	234	E2	11	134
E7	8	6	E2	10	92
E7	10	6	E6	8	141
L2	9	364	E6	9	141
L2	10	364	E4	10	82
L2	10	418		11	82
L2	8	165	E2	8	145
L2	10	165	E2	10	145
L2	8	379	E1	8	237
L2	9	379	E6	8	61
L2	10	379	L2	9	349
L1	10	27	E6	8	82
L2	9	185	E1	8	262
L2	11	185	E1	10	262
L2	9	146	E1	11	380
E1	8	565	E6	9	85
L1	8	344	E6	8	46
L1	10	327	E6	9	46
E1	11	238	L1	9	385
L1	8	20	L1	10	385
L1	9	20	E5A	8	81
L1	11	20	E5A	9	81
E5A	8	25	E4	11	22
E5A	9	25	E6	10	105
E5A	10	25	E1	10	86
E5A	11	25	L2	9	435
L1	8	329	E1	8	579
L1	10	329	E1	10	579
L1	11	329	E5A	9	54
E2	8	349	E5A	,	54

E5A	10	54
L2	8	371
L2	9	371
L2	10	371
E1	9	532
E1	10	532
Ll	10	358
E1	11	536
L2	9	18
· L1	8	65
L1	9	65
L1	10	65
E2	8	159
E2	11	.159
L1	10	350
E2	8	214
E2	9	214
E2	10	214
E5A	8	43
E5A	9	43
E5A	10	43
E5A	11	43
E5B	8	62
E5B	10	62
E1	8	402
E4	8	16
E4	9	16
E4	10	16
E4	9	9
E4	10	9
E4	11	9
E2	8	168
L1	9	287
L1	10	287
L2	8	71
L1	8	10
L1	11	10
E2	9	138
E2	11	138
L1	8	415
E1	8	91
El	9	91
E1	11	91
E5B	8	57
L1	11	26
E2	9	131

_	_			E2	10	217
2	3	4		L1	9	22
E5	8	9		L1	11	22
E5	9	9		L2	9	22
E5	10	9		L2	10	22
E6	9	62				
L1.	9	235		L2	10	13
L2	10	328		L2	11	
L2	11	328		E1	9	525
L2	9	339		E1	11	525
L2	11	339		E6	10	10
E4	8	12		E5	8	11
E4	10	12		E5	10	11
L2	8	86		E5	11	11
L2	11	86		E1	11	77
E2	8	282		E5	8	25
E5	8	10		E5	9	25
E5	9	10		E5	10	25
E5	11	10		E5	11	25
E6	11	83		E1	10	181
E2	8	3		E1	10	101
E2	8	72		L1	9	43
E2	11	72		E5	8	37
L2	10	111		E5	11	37
L2	11	111		E5	8	26
E1	11	112		Ll	8	484
L2	8	139		E1	8	601
L2	11	139		E6	11	64
E1	10	407		E1	11	234
L1	8	421		E1	11	406
L2	8	80		E5	10	46
L2	8	285		E5	11	46
L2	9	285		L1	8	342
E1	9	22		L1	9	342
E1	11	22		L1	11	342
E1	8	475		E1	10	473
E1	8	65		E5	8	27 27
E1	10	65		E5 E5	9 10	27
L1	11	81		E5	11	27
L2	9	417		E2	8	35
L2	10	417		E2	9	35
L2	8	227		E6	8	67
L2	11	227		E6	9	137
E1	10	554 554		E2	9	295
E1	11			E1	8	488
E6	9	37 37		E1	11	488
E6	11 8	24		L1	9	154
E4 E4	9	24		E2	8	11
E4	11	24		E7	9	71
E1	9	319		E1	9	14
L1	9	204		E1	10	14
E1	10	63		E1	11	14
PI RI	8	313		E2	9	227
L1	10	301		E1	9	289
	9	13		E1	10	289
E4				E1	11	289
E4	11	13 45		E2	8	251
E7	10	45		E2	11	251
E7	11 10	45 381		E5	10	73
E1				E6	8	31
E2	8	217		_0	-	

			rica-Az Supermou	repud	28	
E6	9	31		E1	8	191
E6	10	31		E1	9	191
E6	11	31		E2	é e	96
E1	9	640		E2	9	96
E1	10	640		L1	8	332
E4	10	73		Li	9	332
E4	11	73		L1	10	332
E2	8	9		E5	8	12
E2	10	9		E5	10	12
E1	10	250		E5	11	12
	10	73		E1	8	534
E1	8	153		L1	10	412
E2 E1	10	516		Li	11	412
		516		L2	9	29
E1	11 11	607		L2	10	29
E1	11	44		L2	11	29
E7		5		L1	9	216
E6	9	5 524		Li	10	216
E1	8	524		L2	11	257
E1	10	24		L2	8	142
L1	9			L2	9	142
E1	8	369		L2	10	142
E1	9	369			8	71
L2	9	277		E1		
L2	10	277		E1	9	71
L2	11	277		E1 L2	9	178 273
E6	11	96		E2	9	2/3
L2	8	191		E1	8	21
L2	9	191		E1	10	21
E7	8	75		E1	10	336
E7	9	75		E1	11	336
E7	10	75		E2	8	324
E1	11	570		E2	9	324
L2	8	321		E2	10	324
L2	9	321		E2	11	324
L2	9	399		E1	11	62
L2	11	399		E2	8	174
L2	10	346		L1	11	300
E1	11	222		L2	8	172
E7	9	81		L2	10	172
E7	10	81		E1	11	180
L1	8	367		E6	8	113
L1	11	367		E6	9	113
E7	8	14		L2	8	331
E7	10	14		L2	9	331
L1	9	209		L2	10	331
L1	11	209		E1	. 8	105
L1	9	439		E1	9	105
E1	11 9	46		E1	11	105
L1		196		E6	10	42
L1	10	196		E2	10	312
L1	10	456		L2	8	333
L1	11	456		E1	8	197
L2	9	41		E1	11	197
L2	11	41		E2	8	17
E6	9	14		E2	10	17
E6	11	14		E1	11	417
L1	10	199		E2	9	74
E1	9	481		E2	10	74
E1	11	481		E1	11	360
E1	11	164		21	11	300

			HLA-A2 Supermotif P	eptides	
E7	11	27	F	1 1	.0 18
E2	10	340			1 18
E2	8	66		.2 8	
E2	10	66	I	2 1	.0 319
E6	9	92			1 319
E6	10	92	F	1 8	252
E6	11	92	·	1 1	.0 252
E2	8	185	I	.1 8	372
E2	9	185	I	1 1	.0 372
E7	10	36	I	2 1	.1 154
E1	8	141		25 8	
E1	9	39		5 9	
E1	10	39			.1 20
L1	8	103		2 9	
L1	10	103		2 9	
E2	11	118			.0 128
L1	8	382		2 8	
L1	9	382		2 9	
E2	8	205		1 8	
E2	9	205			.1 327
E2	11	205		5 8	
L2	11	119		5 9	
E5	8	2			.0 23
E5	9	2			.1 23
E5	. 10	. 5			.1 197
E5	11	2		5 8	
E6	8	61		5 9	
E6	10	61			.0 40
Ll	11	207			.1 40
L1	8	80		2 8	
L1	9	253			.1 61
L1	11	253		1 8	
L2	9	438			0 203
L2	11	438		2 9	
E5	8	24			.0 313
E5	9	24		1 8	
E5	10	24			.0 319
E5	11	24			1 319
E1	8	50		1 9	
L1	8	370			
L1	9	370		1 8	
L1	10	370			1 156
E1	8	454			0 101
E1	10	454		2 8	
L2	9	424		7 8	
E1	10	494		5 9	
L1	11	464			.0 29
E1 E1	8 11	393 393		2 8	
E2	10	345		2 9	
E2	11	345			.0 .55
E1	10	446		1 8	
E1	9	457		2 9	
L2	8	238			.0 50
L2	9	238		1 9	
L2	10	238			0 401
E5	9	16		2 8	
E5	10	16			0 222
E5	11	16		1 9	
L2	11	275			1 145
		275	_		

			IILA-AZ 3	apennoni i epnae		
E1	8	273		E2	8	78
L1	8	137		E2	9	309
L1	10	137		L1	9	325
E1	10	160		L1	10	325
E2	8	162		E5	9	7
L2	8	309		E5	11	7
L1	8	107		E1	9	305
L2	8	299		E1	10	449
L2	8	24		E2	10	273
L2	11	24		E2	11	273
E5	8	40		E6	8	25
E1	10	6		L1	8	388
L2	10	59		E4	10	36
L1	10	408		L1	8	290
E1	11	296		L2	8	36
E7	9	85		L2	9	36
E7	11	85		L2	11	148
L2	8	408		E4	8	64
L2	10	408		E4	11	64
L2	11	408		L2	9	188
E1	8	467		L2	11	188
E1	9	467		L1	10	362
E1	10	467		E5	8	8
E1	11	467		E5	9	8
Ll	9	169		E5	10	8
E4	8	99		E5	11	8
	10	99		E1	8	248
E4 L1	10	223		E1	9	248
	11	262		L2	11	39
L1 E1	9	173		E2	8	208
E1	10	173		E2	10	208
E2	10	88		E5	8	22
L2	9	50		E5	10	22
L2	11	50		E5	11	22
	9	111		E1	9	474
L1	10	111		E4	8	5
L1	10	316		E4	9	5
E1	10	232		E4	11	5
E2	8	113		E5	8	34
L1		189		E5	9	34
E1	8	189		E5	11	34
E1	10 11	189		L2	9	112
E1	9	415		L2	10	112
E1 L1	8	35		L2	8	278
	9	35		L2	9	278
L1		35		L2	10	278
L1	10	53		L2	10	140
E2	10			L2	11	140
E2	11	53		E1	10	195
E6	9	119		E6	8	120
E6	11	119		E6	10	120
L2	8	176		L2	8	177
L2	9	176		L2	9	177
L2	10	176		E5	8	35
E2	8	29		E5	10	35
E2	10	29		E6	10	97
E1	8	264		L2	9	43
E1	11	264		E5	9	17
E1	9	55		E5	10	17
E1	11	55		E5	8	28
E2	11	136		ca	0	28

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E5	9	28				8	420
E5	10	28				8	42
E1	9	408			L1	8	433
E1	11	408			E2	8	112
E2	9	30			E2	8	47
E6	9	29			E2	10	47
E6	10	29			L1	8	149
E6	11	29			L1	11	149
L1	9	192			E1	9	424
L2	9	165			E1	11	424
E1	8	313				8	53
E1	10	265			E6	9	18
E1	11	265			E6	10	18
L2	8	400				10	84
L2	10	400				11 11	84 165
L2	11	81					518
L2	8	425			E1 E1	9	518
L2	11	425			E7	10	39
L1	8	377				11	39
L1	9	377				11	433
L1	11	377			L2		33
E1	8	56			L2	8 11	33
E1	10	56			E2	9	358
E1	11	56			E2	10	358
E1	11	341			E1	9	121
L2	9	184			E6	8	99
L2	11	184			E1	8	283
L2	8	286 130			E1	9	283
L2	10	130			E1	10	283
L2		130			L1	10	53
L2	11	188			L1	9	61
L1 L1	10 11	188			E2	8	147
E4	9	92			E2	11	147
E4	10	92				9	19
E4	11	92			Ll	11	71
E1	8	23			E6	11	52
El	10	23			E1	8	351
E7	11	11			E1	9	351
E5	11	31			E1	11	351
E1	10	443			E2	8	82
E2	11	286			E4	9	23
L2	9	103			E4	10	23
L2	10	103			E2	9	313
L2	11	103			Ll	10	42
E2	10	45			L2	8	106
L1	9	312			L2	10	106
L2	10	21			E1	9	255
L2	11	21			E1	11	255
E6	11	73			E1	11	307
E2	10	350			E5	9	33
E1	8	312			E5	10	33
E1	9	312			E1	8	557
E1	8	254			E1	9	557
E1	10	254			El	10	557
E6	9	116			E1	8	491
E1	9	460			E1	9	491
E6	11	128			E1	10	491
E1	9	357			E5	8	16
E1	9	114			E5	10	16

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E5	11	16		E5	8	65
E6	11	101		E5	10	65.
E1	10	223		L1	10	368
E1	11	223		Ll	11	368
L2	8	178		L2	8	260
E5	9	36		L2	11	260
E5	11	36		E2	8	289
L1	11	187		E6	11	27
L1	11	41		L1	11	310
	11	521		E1	9	511
E1		540		E1	10	511
E1	9	15		E1	11	511
E2	10			E7	9	15
E7	8	83				
E7	11	83		L2	10	228
E2	8	42		L1	9	466
E2	10	42		L1	11	466
E4	8	18		L1	8	210
E5	10	32		L1	10	210
E1	10	198		E2	8	56
E1	11	198		E2	9	56
E7	8	82		E7	11	55
E7	9	82		E5	10	4
E5	8	31		E1	8	514
E5	11	31		E1	9	514
E5	8	30		E1	11	132
E5	9	30		E1	8	358
E5	11	59		E4	11	90
E5	8	55		E5	10	70
E5	9	55		E1	9	555
E5	11	55		E1	10	555
E5	9	51		E1	11	555
ES	10	51		E5	8	49
ES	11	51		E5	9	49
E1	9	298		E5	11	49
	10	298		E1	8	268
E1		298		E1	9	268
E1	11			E1	10	268
L1	8	119		E1	11	268
L1	10	465				48
E5	11	69		E7	8	
E5	10	60		E7	9	48
E5	11	60		L2	8	368
E1	9	276		E6	8	38
E1	9	563		E6	10	38
E5	8	56		E4	9	69
E5	10	56		E5	9	61
E5	8	52		E5	10	61
E5	9	52		E5	11	61
E5	10	52		E1	8	115
E5	11	52		E2	11	62
E4	10	45		L2	8	337
E4	11	45		L2	11	337
E7	8	79		L1	11	273
E7	9	79		E1	8	277
E7	11	79		E5	9	47
E2	8	139		E5	10	47
E2	10	139		E5	11	47
E2	11	139		L1	8	296
E2	8	94		E7	8	5
		94		E7	9	5
E2	10			E7	11	5
E2	11	94				,

				ILIT-NE C	apermon	repudes		
E1	8	564				E6	8	131
L2	8	245				E6	9	75
E7	10	67				E6	10	75
L1	8	95				Ē2	8	248
L1	10	95				E2	11	248
L1	10	234				L2	10	373.
E4	8	11				L2	11	373
E4	9	11				E5	9	54
E4	11	11				L2	8	381
L1	11	384				E1	10	97
E1	8	306				L1	9	143
E5	11	3				L1	11	143
E1	10	398				E2	9 .	127
E1	11	398				E2	11	127
E2	8	75				E7	9	64
E2	9	75				L2	8	85
E2	11	75				L2	9	85
L1	11	339				L2	10	236
E1	10	47				L2	11	236
E1	11	47				E5	9	78
L1	8	197				E5	10	78
L1	9	197				L2	8	138
E1	9	19				L2	9	138
Ē1	10	19				L2	8	79
L1	8	155				L2	9	79
E1	11	274				L2	9	284
E5	8	1				L2	10	284
E1	10	361				L2	8	416
E4	10	1				L2	10	416
E4	11	1				L2	11	416
L2	8	114				E2	8	216
L2	11	114				E2	9	216
E2	9	71				E2	11	216
E6	8	36				E2	8	196
E6	10	36				E2	11	196
L2	8	269				L1	8	483
L2	10	269				L1	9	483
L2	11	269				E4	8	4
E1	10	389				E4	9	4
E1	9	329				E4	10	4
E1	11	100				L2	10	354
E1	9	600				L2	11	354
E1	8	270				E1	8	94
E1	9	270				E1 E1	9 11	94 442
E1	10	270						
E1	11	270				E6	8 11	110
E1	8	451				E6 L1	8	110 218
E1	11	451				L1	8	184
L1	11	31				L1	9	184
E1	8	300				L2	8	447
E1	9	300				L2	9	447
E2	8	254				L2	8	157
E2	9	254				L2		
E7	8	88				L1	9 10	157 161
L1	10	446				L1	8	459
E1	8	539				L2	9	72
E1	10	539				L2	10	72
E6	9	21				L2	11	72
E1	11 8	397				L1	11	109
L1	0	238						.00

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L1	9	118		E5	8	19
E4	8	30		E5	10	19
E1	8	562		E5	11	19
E1	10	562		L2	8	251
L2	11	211		L2	10	251
E4	11	44		L2	8	327
E5	8	64		L2	11	327
E5	9	64		E1	10	636
E5	11	64		E1	11	636
L2	10	385		L1	9	420
E1	8	258		E1	9	399
E1	11	258		E1	10	399
E1	8	513		E1	11	399
E1	9	513		E1	9	64
E1	10	513		E1	11	64
E7	8	47		E5	8	7
E7	9	47		E5	9	7
E7	10	47		E5	10	7
E4	10	68		E5	11	7
L2	9	336		L2	8	42
E1	10	545		L2	10	42
E1	11	545		E6	10	28
L2	11	167		E6	11	28
L2	11	242		E1	10	31
L2	8	419		E6	8	15
E2	10	353		E6	10	15
L2	8	182		L1	8	152
L2	11	182		L1	11 9	152 373
L2	8	123		L1 L1		373
L2	9	123		E7	11 10	28
E4	8	76		L1	9	302
E4 L2	11 10	76 206		E2	11	14
		90		E7	9	78
L1 L1	9 11	90		E7	10	78
E2	9	211		E2	9 .	288
E2	10	211		L2	8	15
E2	11	211		L2	9	15
E6	11	87		L2	11	15
E2	8	222		E4	8	14
L2	9	95		E4	10	14
L2	10	95		E4	11	14
L2	8	170		L1	11	233
L2	9	170		L1	11	251
L2	10	170		E2	9	48
L2	11	422		E2	11	48
E7	10	20		E2	8	76
L2	8	90		E2	10	76
L1	8	267		E2	9	343
L2	11	358		L2	9	229
E5	8	5 .		L2	11	229
E5	9	5		L2	8	18
E5	10	5		L1	9	211
E5	11	5		L1	11	211
L1	8	16		L1	10	150
L1	10	16		E2	9	218
L2	8	413		E2	11	218
L2	11	413		E1	8	344
L2	9	324		E1	9 · 8	344
L2	11	324		L2	0	230

			HLA-A2 Supermotit	Peptides		
L2	10	230		E1	9	324
L2	8	232		E1	10	324
L2	10	232		E1	11	324
E2	8	57		E1	9	293
E2	11	57		L1	8	29
E1	8	391		L1	8	473
E1	10	391		E1	8	231
L2	8	214		E1	9	231
L2	10	214		E1	10	231
L1	9	260		L2	8	220
L2	8	226		L2	9	220
L2	9	226		L1	11	141
E1		553 -		E2	9	281
E1	8	318		E2	9	225
E1		318		E2	11	225
E2	10	240		L1	8	380
E2	11	240		L1	10	380
L2	8	4		L1	11	380
L2	10	4		L2	8	110
L2	11	4		L2	11	110
L2	10	10		E2	8	234
E5	9	36		L2	10	180
L2	8	297		L1	10	475
L2	10	297		L1	11	475
L2	8	315		L2	8	12
L2	11	315		L2	11	12
L2	9	445		E6	8	9
L2	10	445		E6	11	9
L2	11	445		E1	9	247
E2	9	7		E1	10	247
E2	10	7		E2	9	207
E1	8	109		E2	11	207
E1	9	109		E2	8	23
E2	10	37		E2	9	23
E1	8	125		E6	8	12
E1	9	125		E6		12
L1	11	242		E1	8	547
E4	11	59		E1	9	547 547
L2	8	280		E1	10	
E2	9	302		E1	11	547
E1	8	616		L2	9	266
E7	8	4		L2 L1	11	266 50
E7	9	4		L1	9	50
E7	10	4		L1	10	50
L2	9	244		E1	11	422
E7	11	66 94		L2	8	387
L1	9			L1	8	376
L1	11	94 69		L1	9	376
E1	9			L1	10	376
E1	10 11	69 69		E5	8	30
E1 E7	8	77		L2	10	102
E7	10	77		L2	11	102
E7	11	77		L1	11	286
	8	342		E1	9	350
E2 E2	10	342		E1	10	350
E2 E1	9	343		E2	9	81
E1	10	343		L1	10	86
L1	8	477		Ll	11	86
L1	9	477		L2	9	48
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L2	11	48	•	E1	9	436
E6	10	23		Ll	8	190
L2	8	105		L1	9	190
L2	9	105		L1	11	190
L2	11	105		L2	9	163
E1	9.	490		L2	11	163
E1	10	490		E2	8	348
E1	11	490		Ll	10	393
L2	9	259		E2	10	40
E4	8	89		E5	8	45
Li	9	295		E5	9	45
E1	9	260		E5	11	45
E1	10	260		L1	10	176
L2	9	406		L1	9	344
L2	10	406		E2	9	198
L2	10	303		E2	11	198
E1	11	635		L2	8	144
L1	10	419		L2	10	144
E1	8	354		L2	8	362
E1	9	354		L2	10	362
L2	9	186		E6	8	40
L2	11	186		E5	8	21
L2	8	376		E5	10	21
L2	9	376		E5	11	21
L2	11	376		L1	8	490
L1	9	489		L1	10	490
L1	11	489		E2	9	220
E2	8	260		E2	10	220
E4	8	95		L2	11	404
E4	9	95		E2	9	354
E4	10	95		E2	11	354
L1	8	336		L2	10	183
L1	10	336		L2	8	129
E2	8	245		L2	9	129
E2	9	245		L2	11	129
E2	11	245		E4	10	91
L2	8	240		E4	11	91
E1	8	185		L1	10	340
E1	11	185		L1	11	340
E6	10	7		E2	10	249
L2	9	135		E5	9	71
L2	11	135		L1	9	409
E4	8	85		E1	8	294
E4	9	85		E1	8	556
E4	10	85		E1	9	556
E2	8	190		E1	10	556
E2	10	190		E1	11	556
E1	11	331		E5	8	15
E2	8	201		E5	9	15
E2	9	201		E5	11	15
L1	11	7		E7	9	7
E2	10	316		E5	8	50
L1	8	282		E5	10	50
L1	10	282		E5	11	50
E2	10	151		E1	10	297
E2	9	257		E1	11	297
E2	10	257		E7	8	86
E2	11	257		E7	10	86
L2	9	152		E2	9	93
E1	8	436		E2	11	93

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L1	8	378	I	55	9	13
L1	10	378	F	35	10	13
L2	9	374	F	35	11	13
L2	10	374	1	.2	10	149
L2	11	374	F	25	9	12
L2	9	326			10	12
L1	8	170			11	12
L2	8	124			8	55
E7	8	49			9	100
E1	11	30			8	212
L1	9	151			9	212
L2	9	14			10	212
L2	10	14			8	290
E1	9	57			9	290
E1	10	57			10	290
E1	8	526			9	224
E1	10	526			11	224
E1	10	117			10	88
L1	9	92			11	88
L1	11	92			10	263
L2	8	364				263
L2	11	364			9	191
E2	9	85			8	345
E2	10	85			10	332
E6	9	39			11	332
E2	8	219				78
E2	10	219			9	82
E2	11	219			10	82
E1	8	232			11	82
E1	9	232				202
E6	8	142				202
E2	8	228 .				223
L2	8-	37				63
E5	8	14				63
E5	9	14				101
E5 '	10	14				329
L2	8	136				329
L2	10	136				8
L2	11	136				338
E4	8	70				239
E5	8 -	62				239
E5	9	62				519 98
E5	10	62				252
E5	11	62				252
E1 L1	11	116 91				96
L1	10	91				96
L1	8	333				96
L1	9	333				291
L1	11	333				291
E4	8	86				291
E4	9	86				283
E4	11	86				21
E4	8	93				25
E4	9	93				25
E4 E4	10	93				21
E4	11	93				21
L2	9	150				21
L2	11	150				192
E5	8	13				192
	-	~~	-			

E1	9	333		ь2	10	325
E1	10	333		L2	9	363
L2	8	30		E2	10	148
L2	9	30		L1	10	328
L2	10	30		E1	11	238
L2	11	30		L1	8	20
L1	8	191		L1	11	20
L1	10	191		E5	8	24
L2	8	164		E5	9	24
L2	10	164		E5	10	24
E7	10	12		Ll	8	330
L1	9	394		L1	10	330 330
E5	11	27		L1	11	
E5	10	32		E7	9	68
E5	11	32		E5	9 10	20 20
E2	9	41		E5	10	72
E2 E4	11	41 17		L1 L1	11	72
	8	17		E4	9	2
E4	9	275		E4	10	2
E1 L1	10	73		E4	11	2
	9 10	73		E2	10	58
L1 E5	8	48		L2	10	120
E5	9	48		L2	11	120
				E5	8	3
E5 E5	10 8	48 46		E5	9	3
E5	10	46		E5	10	3
E5	11	46		E5	11	3
E4	8	10	-	L2	10	198
E4	9	10		E6	10	53
E4	10	10		E6	11	53
Li1	8	383		E2	8	132
E2	8	128		E5	8	41
E2	10	128		E5	9	41
E1	9	79		E5	10	41
E1	- 9	444		E5	11	41
L2	10	359		L1	8	97
L2	11	359		E2	11	320
E5	8	6		E1	9	426
E5	9	6		E1	8	340
E5	10	6		E5	8	14
E5	11	6		E5	9	14
E2	10	287		E5	11	14
L1	9	177		E5	8	18
L1	11	177		E5	9	18
L2	8	394		E5	10	18
L2	11	394		E5	11	18
E4	8	83		E1	11	464
E4	9	83		E5	8	58
E4	10	83		E1	8	510
E4	11	83		E1	10	510
L2	9	231		El	11	510
L2	11	231		E1	8	267
E1	9	362		E1	9	267
L2	9	233		E1	10	267
E7	8	6		E1	11	267
E7	10	6		E2	10	92
L2	9	145		E6	8	141
L2	10	414		E6	9	141
L2	8	325		E4	10	81

Table VIIIC HPV11 HLA-A2 Supermotif Peptides

E4	11	81
E1	8	530
E1	11	530
E2	8	145
E2	10	145
E1	8	237
E6	8	82
L2	8	348
El	8	262
E1	10	262
		85
E6	9	
E1	11	380
E6	8	44
E6	10	44
E6	11	44
E6	8	46
E6	9	46
E5	9	81
Li	9	386
L1	10	386
L2	11	70
E4	10	22
E4	11	22
E6	10	105
E1	10	86
L2	9	431
E1	8	579
E1	10	579
		54
€5	8	
E5	9	54
E5	10	54
E2	9	138
E2	11	138
L1	9	246
L2	8	367
L2	9	367
E1	9	532
Ε1	10	532
L1	10	359
E1	11	536
E5	11	61
	9	17
L2		
L1	8	65
Ll		65
L1	10	65
E2	8	159
E2	11	159
L1	10	351
E2	8	214
E2	10	214
E2	11	214
	8	305
L2	10	305
E5		43
E5		43
E5		43
E5	11	43
Li	9	288
61	10	288
E1		402
	-	

L1	11	26
E4	8	16
E4	9	16
E4	10	16
E4	9	9
E4	10	9
E4	11	9
E2	8	168
E1	10	502
L1	8	10
L1	11	10
L1	8	416
E1	8	91
E1	9	91
E1	11	91
E2	9	131

SF 1168095 v1

Table IX HLA-A3 Supermotif Peptides

				•	•			
1	2	3	4		HPV16	E1	10	278
HPV16	E1	8	316		HPV16	E1	9	544
HPV16	E1	8	205		HPV16	E1	8	306
HPV16	E1	9	112		HPV16	E1	9	305
HPV16	E1	9	69		HPV16	E1	8	454
HPV16	E1	11	459		HPV16	E1	9	454
HPV16	E1	8	406		HPV16	E1	8	420
HPV16	E1	9	406		HPV16	E1	10	420
HPV16	E1	8	82		HPV16	E1	8	422
HPV16	E1	9	405		HPV16	E1	11	422
HPV16	E1	10	405		HPV16	E1	8	273
HPV16	E1	10	114		HPV16	E1	10	273
HPV16	E1	11	114		HPV16	E1	9	202
HPV16	E1	8	304		HPV16	E1	11	202
HPV16	E1	10	304		HPV16 HPV16	E1 E1	9	567 543
HPV16	E1	9	101		HPV16	E1	10	543
HPV16	E1	11	101		HPV16	E1	9	386
HPV16	E1	8	81		HPV16	E1	8	396
HPV16		9	81 368		HPV16	E1	9	196
HPV16		11	573		HPV16	E1	10	190
HPV16	E1	11	384		HPV16	E1	10	302
	E1	8	335		HPV16	E1	8	245
HPV16		11	548		HPV16	E1	8	600
HPV16		8	603		HPV16	E1	11	600
HPV16		10	221		HPV16	E1	8	143
HPV16	E1	9	288		HPV16	E1	9	419
HPV16	E1	11	140		HPV16	E1	11	419
HPV16	E1	9	392		HPV16	E1	8	118
HPV16		8	463		HPV16	E1	9	109
HPV16	E1	9	453		HPV16	E1	10	619
HPV16	E1	10	453		HPV16	E1	11	313
HPV16	E1	9	219		HPV16	E1	9	432
HPV16	E1	10	71		HPV16	E1	11	390
HPV16		11	242		HPV16	E1	9	484
HPV16	E1	9	272		HPV16	E1	8	621
HPV16	E1	11	272		HPV16	E1	9	421
	E1	10	174		HPV16	E1	10	314
HPV16	E1	10	496		HPV16	E1	9	497
HPV16		9	216		HPV16	E1	9	315 72
HPV16	E1	10	68		HPV16 HPV16	E1	8	289
HPV16	E1	11	473		HPV16	E1 E1	8	407
HPV16	E1	11	194 369		HPV16	E1	11	407
HPV16	E1 E1	10 10	401		HPV16	E1	11	200
HPV16	E1	9	204		HPV16	E1	11	565
HPV16	E1	10	111		HPV16	E1	8	498
HPV16	E1	11	400		HPV16	E1	8	197
HPV16	E1	10	610		HPV16	E1	8	275
HPV16	E1	10	483		HPV16	E1	11	275
HPV16	E1	10	394		HPV16	E1	8	217
HPV16	E1	10	276		HPV16	E1	11	217
HPV16	E1	9	277		HPV16	E1	8	545
HPV16		11	277		HPV16	E1	9	274
HPV16	E1	10	474		HPV16	E1	8	425
HPV16	E1	9	620		HPV16	E1	9	509
HPV16	E1	9	191		HPV16	E1	8	20
HPV16	E1	10	243		HPV16	E1	9	20
HPV16	E1	9	222		HPV16	E2	8.	40
HPV16	E1	8	278		HPV16	E2	8	300

Table IX HLA-A3 Supermotif Peptides

				•			
HPV16	E2	9	174	HPV16	E5	8	51
HPV16	E2	9	294	HPV16	E5	9	22
HPV16	E2	11	294	HPV16	E5	11	48
HPV16	E2	10	25	HPV16	E5	10	70
HPV16	E2	10	246	HPV16	E5	10	21
HPV16	E2	8	233	HPV16	E5	9	50
HPV16	E2	10	233	HPV16	E6	9	7
HPV16	E2	9	204	HPV16	E6	11	7
HPV16	E2	9	346	HPV16	E6	8	68
	E2	10	168	HPV16	E6	9	143
HPV16		10	163	HPV16	E6	11	143
	E2	10	156	HPV16	E6	10	37
HPV16		11	230	HPV16	E6	11	37
	E2	9	29	HPV16	E6	10	32
	E2	10	290	HPV16	E6	11	105
	E2	11	35	HPV16	E6	8	48
	E2	8	252	HPV16	E6	11	52 92
	E2	10	267	HPV16 HPV16	E6	10 9	33
	E2	8	45	HPV16	E6	8	34
HPV16		11	215 347	HPV16	E6	9	107
	E2	9	268	HPV16	E6	10	106
	E2 E2	11	268	HPV16	E6	8	144
	E2 E2	9	103	HPV16	E6	10	144
HPV16		10	103	HPV16	E6	11	144
	E2	9	335	HPV16	E6	9	134
	E2	11	282	HPV16	E6	8	102
	E2	8	84	HPV16	E6	9	116
	E2	9	296	HPV16	E6	11	5
	E2	11	296	HPV16	E6	10	6
	E2	9	284	HPV16	E6	8	94
	E2	11	266	HPV16	E6	9	93
	E2	9	60	HPV16	E6	10	139
HPV16	E2	8	235	HPV16	E6	9	67
HPV16	E2	10	57	HPV16	E6	8	77
HPV16	E2	9	37	HPV16	E7	10	68
HPV16	E2	11	37	HPV16	E7	10	88
HPV16	E2	8	7	HPV16	E7	9	89
HPV16	E2	8	165	HPV16	E7	8	53
HPV16	E2	11	317	HPV16	E7	9	41
HPV16	E2	8	269	HPV16		8	70
	E2	10	269		L1	11	372
	E2	8	104	HPV16		9	162
	E2	9	104	HPV16	L1	10	373 233
	E2	11	81	HPV16	L1 L1	11	70
	E2	8	61	HPV16	Ll	11	70
	E2	8	297	HPV16	L1	8	128
	E2	10	297	HPV16	L1	8	249
	E2	11	297	HPV16	L1	9	484
	E2	10	334	HPV16	Ll	10	484
	E2 E2	8	285	HPV16	LI	10	397
	E2 E2	11	333	HPV16	L1	8	270
	E2 E2	9	58	HPV16	Ll	9	270
	E2 E2	11	58	HPV16	Ll	11	113
	E2	9	321	HPV16	Ll	10	378
	E2	10	102	HPV16	Ll	8	494
	E2	11	102	HPV16	L1	10	494
	E5	11	20	HPV16	L1	8	236
HPV16		8	72	HPV16	L1	8	282
HEVIO.							

Table IX HLA-A3 Supermotif Peptides

				HLA-A3 Supermotti i	repudes			
HPV16		11	446		HPV16	T.1	11	520
HPV16		9	356		HPV16		8	522
HPV16		10	142		HPV16		9	522
HPV16		8	93		HPV16		10	516
HPV16		8	438		HPV16		11	516
HPV16		9	143		HPV16		9	3.79
HPV16		9	374		HPV16		11	36
HPV16		10	501		HPV16		10	91
			501		HPV16		9	48
HPV16		11	90		HPV16		10	326
		11	90		HPV16		10	447
HPV16			46		HPV16		8	357
HPV16		11 11	69		HPV16		10	47
HPV16		9	495		HPV16		10	126
HPV16		11	495		HPV16		10	161
HPV16		11	87		HPV16		9	38
HPV16		11	325		HPV16		10	275
HPV16		10	58		HPV16		9	470
HPV16		9	383		HPV16		11	470
HPV16		9	296		HPV16		10	288
HPV16		9	460		HPV16		11	288
HPV16		10	460		HPV16		10	293
HPV16		8	190		HPV16		8	13
HPV16		9	77		HPV16		11	13
HPV16		10	247		HPV16		9	82
HPV16		11	515		HPV16		9	15
HPV16		9	497		HPV16		9	31
HPV16		11	331		HPV16		9	283
HPV16		8	181		HPV16		11	59
HPV16		11	354		HPV16		10	300
HPV16		10	280		HPV16		11	226
HPV16		10	179		HPV16		10	26
HPV16		9	100		HPV16		9	61
HPV16		11	482		IPV16		8	32
HPV16		10	253	1	IPV16	L2	9	294
HPV16		8	271	1	IPV16	L2	8	454
HPV16		8	518		HPV16	L2	9	240
HPV16		9	518	1	HPV16	L2	11	292
HPV16		10	518	1	HPV16	L2	10	215
HPV16		11	518	1	HPV16	L2	8	450
HPV16		8	49	1	HPV16	L2	9	450
HPV16		8	375	I	HPV16	L2	10	450
HPV16	L1	8	519	1	HPV16	L2	11	450
HPV16	L1	9	519	1	HPV16	L2	11	80
HPV16	L1	10	519	1	HPV16	L2	10	221
HPV16	L1	11	519		HPV16		9	310
HPV16	L1	В	521		HPV16		9	12
HPV16	L1	9	521		HPV16		9	305
HPV16	L1	10	521		HPV16		11	305
HPV16	Ll	8	523		HPV16		8	5
HPV16	L1	9	327		HPV16		9	315
HPV16	L1	10	114		HPV16		8	298
HPV16	L1	11	252		HPV16		10	69
HPV16	L1	9	448		HPV16		11	313
HPV16	L1	9	517		HPV16		10	14
HPV16		10	517		HPV16		9	212
HPV16	L1	11	517		HPV16		8	213
HPV16	L1	8	520		HPV16		10	81
HPV16		9	520		IPV16		8	311
HPV16	L1	10	520	I	HPV16	L2	8	295

Table IX HLA-A3 Supermotif Peptides

HPV16	L2	11	295		HPV18	E1	9	647
HPV16	L2	10	211		HPV18	E1	9	468
HPV16	L2	11	287		HPV18	E1	10	468
HPV16	L2	9	222		HPV18	E1	10	401
HPV16	L2	11	210		HPV18	E1	8	292
HPV16	L2	10	447		HPV18	E1	10	283
HPV16	L2	11	447		HPV18	E1	9	281
HPV16	L2	8	453		HPV18	E1	8	313
HPV16	L2	9	453		HPV18	E1	8	285
HPV16	L2	11	303		HPV18	E1	10	285
HPV16	L2	9	228		HPV18	E1	10	570
HPV18	E1	11	397		HPV18		8	224
HPV18	E1	11	546		HPV18	E1	11	224
HPV18	E1	11	466		HPV18	E1	9	571
HPV18	E1	9	284		HPV18		11	480
HPV18	E1	11	284		HPV18	E1	9	229
HPV18	E1	8	413		HPV18		9	312
HPV18	E1	9	413		HPV18		8	429
HPV18	E1	9	412		HPV18		11	429
HPV18	E1	10	412		HPV18		9	574
HPV18	E1	8	311		HPV18		9	428
HPV18	E1	10	311		HPV18		8	119
HPV18		11	437		HPV18		9	119
HPV18	E1	11	196		HPV18		10	119
HPV18		9	78		HPV18		9	393
HPV18		10	78		HPV18		9	551
HPV18		11	78		HPV18		8	252
HPV18		8	203		HPV18		8	607
HPV18		10	228			E1	11	607
HPV18		11	391		HPV18		11	200
HPV18		11	637		HPV18		9	426
HPV18		8	342		HPV18		11	426 80
HPV18		8	610		HPV18 HPV18		9	80
HPV18		9	115		HPV18		11	102
HPV18		10	115		HPV18		11	320
HPV18		10	309		HPV18		8	117
HPV18		9	104		HPV18		10	117
HPV18		9	460		HPV18		11	117
HPV18		10	463 470		HPV18		10	321
HPV18		9	399		HPV18		10	93
HPV18		9	226		HPV18		9	322
HPV18		8	465		HPV18		10	197
HPV18		8	212		HPV18		8	414
HPV18		9	223		HPV18		11	414
HPV18		11	92		HPV18		8	572
HPV18		9	279		HPV18		11	572
HPV18		11	279		HPV18	E1	8	323
HPV18		11	249		HPV18	E1	8	81
HPV18		8	270		HPV18	E1	8	280
HPV18		9	198		HPV18	E1	10	280
HPV18	E1	8	282		HPV18	E1	11	339
HPV18		11	282		HPV18	E1	8	432
HPV18		11	569		HPV18	E1	9	516
HPV18		8	552		HPV18	E1	9	536
HPV18	E1	8	116		HPV18	E1	10	268
HPV18	E1	9	116		HPV18	E1	10	408
HPV18	E1	11	116		HPV18		8	19
HPV18	E1	8	461		HPV18		9	19
HPV18	E1	9	439		HPV18	E2	9	269

Table IX HLA-A3 Supermotif Peptides

HPV18	F2	10	269	1	HPV18	E2	10	157
HPV18		11	269		HPV18		9	335
HPV18		9	82	i	HPV18	E2	11	335
HPV18		11	82		HPV18		9	62
HPV18		8	270		HPV18		11	62
HPV18	E2	9	270	. 1	HPV18	E2	8	322
HPV18		10	270	1	HPV18	E2	10	173
HPV18		11	270	1	HPV18	E2	10	143
HPV18		8	301	1	HPV18	E2	11	228
HPV18	E2	11	156	1	HPV18	E6	9	68
HPV18		11	31	1	HPV18	E6	10	27
HPV18	E2	10	210	1	HPV18	E6	10	58
HPV18	E2	10	268	1	HPV18	E6	10	83
HPV18	E2	11	268	1	EPV18	E6	8	29
HPV18	E2	8	85		HPV18		11	40
HPV18	E2	10	291		HPV18		8	43
HPV18	E2	8	338		HPV18		11	47
HPV18	E2	10	19		IPV18		8	97 .
HPV18		8	289		IPV18		11	97
HPV18		8	68		IPV18		8	139
HPV18		11	18		HPV18		11	139
HPV18		9	152		IPV18		8	117
HPV18		11	238		IPV18		9	117
HPV18		11	8		HPV18		10 9	117
HPV18		11	333 81		IPV18		10	101
HPV18		10 9	144		IPV18		10	41
HPV18 HPV18		8	44		IPV18		9	1
HPV18		9	67		IPV18		10	î
HPV18		9	297		IPV18		8	100
HPV18			297		IPV18		11	100
HPV18		9	107		IPV18		10	95
HPV18		10	107	I	IPV18	E6	11	114
HPV18	E2	8	170	I	IPV18	E6	9	111
HPV18	E2	9	285	I	IPV18	E6	9	144
HPV18	E2	9	64	1	IPV18	E6	10	144
HPV18	E2	9	288		IPV18		11	144
HPV18		8	272		IPV18		9	59
HPV18	E2	9	272		IPV18		9	84
HPV18		9	33		IPV18		8	72
HPV18		11	80		IPV18		9	63
HPV18		10	2		(PV18		11	63
HPV18		11	119		IPV18		8	77 43
HPV18		10	61		IPV18		10 11	43
HPV18 HPV18		8 10	122 305		IPV18		11	48
		8	11		IPV18		9	59
HPV18 HPV18		8	298		IPV18		11	74
HPV18		10	298		IPV18		11	61
HPV18		11	298		IPV18		9	50
HPV18		10	229		IPV18		8	60
HPV18		9	230		IPV18		10	75
HPV18		8	153		IPV18		11	195
HPV18		8	286	F	IPV18	L1	8	225
HPV18		11	286		IPV18		11	268
HPV18		10	120		IPV18		9	419
HPV18	E2	9	211		IPV18		10	196
HPV18		8	231		IPV18		9	552
HPV18		10	334		IPV18		10	552
HPV18	E2	8	212	F	[PV18	r1	8	163

Table IX HLA-A3 Supermotif Peptides

HPV18	L1	11	222	HPV18		9	555
HPV18	L1	10	310	HPV18		11	555
HPV18	L1	8	493	HPV18		8	485
HPV18	L1	10	418	HPV18		9	362
HPV18	L1	8	284	HPV18		11	92
HPV18	L1 .	11	122	HPV18		10	149
HPV18	L1	9	520	HPV18		8	474
HPV18		10	520	HPV18		9	197
HPV18		8	305	HPV18		8	554
HPV18		9	305	HPV18		10	554
HPV18		11	148	HPV18		9	473
HPV18		10	330	HPV18		8	553
HPV18		11	203	HPV18		9	553
HPV18		8	317	HPV18		11	553
HPV18		8	59	HPV18		8	105
HPV18		8	530	HPV18		9	331
HPV18		9	530	HPV18		11	71 126
HPV18		10	530	HPV18		10	
HPV18		8	271	HPV18 HPV18		10 10	361 161
HPV18		11	482	HPV18		8	230
HPV18		11	535	HPV18		10	230
HPV18		10	177	HPV18		9	73
HPV18		11	360 505	HPV18		10	286
HPV18		10	125	HPV18		8	12
HPV18		8 11	125	HPV18		11	12
HPV18 HPV18		10	103	HPV18		11	354
HPV18		9	178	HPV18		9	273
HPV18		9	104	HPV18		11	109
HPV18		8	531	HPV18		9	260
HPV18		9	531	HPV18	L2	8	443
HPV18			496	HPV18	L2	9	276
HPV18		9	224	HPV18	L2	11	306
HPV18		8	558	HPV18	L2	11	58
HPV18		10	558	HPV18	L2	10	25
HPV18	L1	11	558	HPV18	L2	9	60
HPV18	L1	10	57	HPV18	L2	11	292
HPV18	L1	10	282	HPV18		10	210
HPV18	L1	10	16	HPV18		11	210
HPV18	L1	8	550	HPV18		10	34
HPV18		11	550	HPV18		9	287
HPV18		8	540	HPV18		8	1
HPV18		10	472	HPV18		9	1
HPV18		10	412	HPV18		10 11	1
HPV18		10	315	HPV18		10	79
HPV18		11	366	HPV18		11	285
HPV18		9	484	HPV18		8	357
HPV18		11	102	HPV18		11	209
HPV18 HPV18		11 9	547 112	HPV18		8	439
HPV18		9	135	HPV18		9	439
		8	561	HPV18		10	439
HPV18 HPV18		10	548	HPV18		11	439
HPV18		10	551	HPV18		10	216
HPV18		11	551	HPV18		11	258
HPV18		9	127	HPV18		9	11
HPV18		10	93	HPV18	L2	11	298
HPV18		9	150	HPV18		10	281
HPV18		11	518	HPV18	L2	11	281
HPV18		8	306	HPV18	L2	9	308
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Table IX HLA-A3 Supermotif Peptides

HPV18	L2	10	364	HPV31 E1	9	157
HPV18	L2	10	68	HPV31 E1	8	386
HPV18		10	220	HPV31 E1	9	386
HPV18	L2	9	211	HPV31 E1	8	225
HPV18	L2	10	211	HPV31 E1	9	196
HPV18		10	110	HPV31 E1	11	222
HPV18		8	212	HPV31 E1	9	78
HPV18		9	212	HPV31 E1	10	78
HPV18		9	365	HPV31 E1	11	78
HPV18		10	235	HPV31 E1	11	162
HPV18		10	13	HPV31 E1	8	478
HPV18		9	111	HPV31 E1	11	453
HPV18		8	288	HPV31 E1	11	174
HPV18		11	288	HPV31 E1	9	268
HPV18		8	261	HPV31 E1	9	544
HPV18		8	366	HPV31 E1	10	381
HPV18		10	293	HPV31 E1	9	184
HPV18		9	217	HPV31 E1	10	110
HPV18		9	80	HPV31 E1	11	380
HPV18		9	221	HPV31 E1	9	441
HPV18		9	236	HPV31 E1	10	441
HPV18		8	2	HPV31 E1	10	590
HPV18		9	2	HPV31 E1	10	374
HPV18		10	2	HPV31 E1	9	412
HPV18		11	234	HPV31 E1	10	454
HPV18		9	14	HPV31 E1	8	286
HPV18		8	81	HPV31 E1	9	202
HPV18		8	112	HPV31 E1	10	543
HPV18		11	436	HPV31 E1	11	542
HPV31		8	296	HPV31 E1	10	256
HPV31		8	185	HPV31 E1	9	437
HPV31		9	111	HPV31 E1	8	258
HPV31		11	439	HPV31 E1	10	258
HPV31		8	81	HPV31 E1	9	285
HPV31		11	370	HPV31 E1	8	255
HPV31		10	263	HPV31 E1	11	255
HPV31		10	113	HPV31 E1	9	257
HPV31		11	113	HPV31 E1	11	257
HPV31		9	477	HPV31 E1	8	400
HPV31		8	284	HPV31 E1	10	400
HPV31		10	284	HPV31 E1	8	253
HPV31		9	100	HPV31 E1	10	253
HPV31		11	100	HPV31 E1	9	547
HPV31		8	620	HPV31 E1	8	601
HPV31		11	364	HPV31 E1	8	117
HPV31		9	366	HPV31 E1	8	376
HPV31		11	528	HPV31 E1	10	170
HPV31		11	348	HPV31 E1	9	524
HPV31		8	80	HPV31 E1	8	580
HPV31		9	80	HPV31 E1	11	580
HPV31		10	201	HPV31 E1	9	399
HPV31		8	583	HPV31 E1	11	399
HPV31		8	315	HPV31 E1	9	176
HPV31		8	443	HPV31 E1	10	267
HPV31		9	372	HPV31 E1	10	599
HPV31		10	436	HPV31 E1	11	293
HPV31		11	566	HPV31 E1	8	438
HPV31		9	433	HPV31 E1	9	401
HPV31		9	252	HPV31 E1	11	98
HPV31		11	252	HPV31 E1	10	294
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Table IX HLA-A3 Supermotif Peptides

				-			
HPV31	R1	11	281	HPV31	E2	9	127
HPV31		9	295	HPV31		8	219
HPV31		8	269	HPV31	E2	9	60
HPV31		8	387	HPV31	E2	10	290
HPV31		11	387	HPV31	32	10	57
HPV31		11	180	HPV31	32	8	238
HPV31	E1	8	545	HPV31	E2	10	238
HPV31		11	545	HPV31	E2	10	25
HPV31	E1.	8	177	HPV31	E2	9	37
HPV31		10	349	HPV31	E2	11	37
HPV31	E1	9	254	HPV31	E2	8	7
HPV31	E1	8	413	HPV31	E2	10	276
HPV31	E1	8	434	HPV31		11	324
HPV31	E1	8	197	HPV31	32	11	216
HPV31	E1	8	525	HPV31		8	104
HPV31	E1	10	223	HPV31		9	104
HPV31	E1	8	405	HPV31		8	81
HPV31	E1	9	489	HPV31		10	341
HPV31		8	19	HPV31		8	128
HPV31	E2	9	277	HPV31		8	292
HPV31	E2	8	278	HPV31		8	240
HPV31	E2	9	229	HPV31		10	146
HPV31	E2	10	229	HPV31		11	340
HPV31	E2	8	61	HPV31		9	147
HPV31	E2	9	291	HPV31		9	58
HPV31	E2	9	239	HPV31		11	58
HPV31	E2	10	228	HPV31		9	328
HPV31	E2	11	228	HPV31		10	102
HPV31	E2	8	307	HPV31		11	102
HPV31	E2	10	307	HPV31		11	20
HPV31	E2	11	145	HPV31		11	48
HPV31	E2	8	40	HPV31		9	22
HPV31	E2	9	301	HPV31		8	51
HPV31	E2	11	301	HPV31			21
HPV31	E2	9	174	HPV31		9	50
HPV31	E2	10	174	HPV31		10	18
HPV31	E2	10	204	HPV31		9	136
HPV31	E2	9	80	HPV31		10	63
HPV31	E2	9	168	HPV31		11	98
HPV31	E2	10	168	HPV31		9	57
HPV31	E2	8	231	HPV31		8	20
HPV31		10	235	HPV31		10	25
HPV31	E2	11	235	HPV31			45
HPV31		9	29	HPV31		9	47
HPV31		11	35	HPV31		8	95
HPV31		10	297	HPV31		10	85
HPV31		10	15	HPV31		8	61
HPV31		11	15	HPV31		8	137
HPV31		8	304	HPV31		9	72
HPV31		11	304	HPV31		9	100
HPV31		11	275	HPV31		10	99
HPV31		9	205	HPV31		9	127
HPV31		11	14	HPV31		9	109
HPV31		11	4	HPV31		8	27
HPV31		9	103	HPV31		11	17
HPV31		10	103	HPV31		9	82
HPV31		9	342	HPV31		8	87
HPV31		11	78	HPV31		9	86
HPV31		9	303	HPV31		8	73
HPV31	E2	10	254	HPV31	46	10	132

Table IX HLA-A3 Supermotif Peptides

HPV31		11	132		HPV31		11	472
HPV31		8	70			L1	11	306
HPV31	E6	11	70			Ll	8	156
HPV31	E6	10	81		HPV31	L1	11	329
HPV31	E7	10	68		HPV31		10	255
HPV31	E7	10	88		HPV31	L1	10	154
HPV31	E7	9	89		HPV31	L1	10	476
HPV31	E7	9	54		HPV31	L1	11	476
HPV31	E7	10	53		HPV31	L1	9	75
HPV31	E7	8	70		HPV31	L1	11	457
HPV31	E7	8	55		HPV31	L1	9	490
HPV31	L1	11	347		HPV31	L1	10	490
HPV31	L1	10	348		HPV31	L1	11	490
HPV31	L1	11	426		HPV31	L1	10	228
HPV31	L1	11	208		HPV31	L1	9	51
HPV31	L1	8	491		HPV31	L1	10	51
HPV31	L1	9	491		HPV31	L1	9	358
HPV31	L1	10	491		HPV31	L1	8	23
HPV31	L1	11	491		HPV31	L1	8	492
HPV31	L1	8	103		HPV31	L1	9	492
HPV31	L1 ·	8	224		HPV31	L1	10	492
HPV31	L1	9	459		HPV31	L1	9	271
HPV31	L1	10	459		HPV31	L1	8	246
HPV31	L1	10	372		HPV31	L1	9	302
HPV31	L1	8	245		HPV31	L1	10	89
HPV31	L1	9	245		HPV31	L1	9	423
HPV31	L1	11	88		HPV31	L1	9	354
HPV31	L1	10	353		HPV31	L1	8	494
HPV31	L1	10	270		HPV31	L1	10	494
HPV31	L1	8	469		HPV31	L1	11	494
HPV31	L1	10	469		HPV31	L1	8	493
HPV31	L1	8	211		HPV31		9	493
HPV31	L1	8	257		HPV31	L1	11	493
HPV31	L1	11	421		HPV31		11	44
HPV31	L1	9	331		HPV31		11	10
HPV31	L1	10	117		HPV31		10	66
HPV31	L1	8	68		HPV31			22
HPV31	L1	10	68		HPV31		10	301
HPV31	L1	8	413		HPV31		10	422
HPV31	L1	9	349		HPV31		8	332
HPV31	L1	9	118		HPV31		11	62
HPV31	L1	10	427		HPV31			21
HPV31		10	357		HPV31		10	101
HPV31		8	431		HPV31		10	136
HPV31		8	65		HPV31		9	12
HPV31		11	65		HPV31			250
HPV31		11	20		HPV31			50
HPV31		9	470		HPV31			50
HPV31		11	470		HPV31			445
HPV31		11	300		HPV31			445
HPV31		10	32		HPV31			281
HPV31		11	227		HPV31			281
HPV31		8	496		HPV31			286
HPV31		9	496		HPV31		11	13
HPV31		8	165		HPV31		9	15
HPV31		10	222		HPV31			276
HPV31		10	489		HPV31			59
	L1	11	489		HPV31 HPV31			221
HPV31		10	411		HPV31		9	61 26
HPV31	L1	9	472		HP V 3 I	LIZ	10	20

Table IX HLA-A3 Supermotif Peptides

HPV31	L2	10	38	HPV33	E1	11	377
HPV31	L2	11	280	HPV33	E1	10	566
HPV31	L2	11	205	HPV33	E1	11	541
HPV31	L2	9	287	HPV33	E1	11	99
HPV31	L2	8	447		E1	9	537
HPV31	L2	11	292	HPV33	E1	10	214
HPV31	L2	11	285	HPV33	E1	8	242
HPV31	L2	9	217	HPV33	E1	10	295
HPV31	L2	10	210	HPV33	E1	8	19
HPV31		8	443	HPV33	E1	9	19
HPV31	L2	9	443	HPV33	E1	10	449
HPV31	L2	10	443	HPV33		8	456
HPV31		11	443		E1	9	385
HPV31		8	235	HPV33			212
HPV31		9	298		E1	9	446
HPV31		11	298	HPV33	E1	10	446
HPV31		9	308	HPV33	E1	8	451
HPV31		8	2 .	HPV33	E1	9	265
HPV31		10	2	HPV33	E1		265
HPV31		11	2	HPV33	E1	8	399
HPV31		8	5		E1	9	399
HPV31		10	69	HPV33	E1	9	209
HPV31		11	306		E1		235
HPV31		10	14	HPV33	E1		480
HPV31		9	207	HPV33	E1		327
HPV31		10	207	HPV33	E1		256
HPV31		8	208	HPV33	E1		573
HPV31		9	208	HPV33			266
HPV31		11	80				266
HPV31		8	40	HPV33	E1		267
HPV31		8	288	HPV33	E1		268
HPV31		11	288				268 400
HPV31		10	206	HPV33			400
HPV31		11	206	HPV33 HPV33	E1 E1		210
HPV31		9	39		E1		210
HPV31		10	293	HPV33			538
HPV31		10	81		E1		187
HPV31		11	232	HPV33	E1		236
HPV31		9	82	HPV33	E1	11	520
HPV31 HPV31		10	440 440	HPV33	E1	10	394
HPV31		11	446	HPV33	E1		197
HPV31		9	446			11	393
HPV31		9	223	HPV33		10	612
HPV31		11	296		E1	11	412
HPV31		8	96			10	603
HPV33		11	383		E1		387
HPV33		11	104	HPV33	E1		425
HPV33		8	596	HPV33	E1		467
HPV33		8	81		E1		271
	E1	8	297		E1	10	271
HPV33		10	297	HPV33	E1		270
HPV33		8	633	HPV33			270
HPV33		11	633	HPV33	E1		269
HPV33	E1	10	276	HPV33	E1	9	215
HPV33	E1	9	490	HPV33	E1	11	466
HPV33		8	614	HPV33	E1	10	413
HPV33		9	78	HPV33	E1	8	481
HPV33		10	78	HPV33	E1	9	298
HPV33	E1	11	78	HPV33	E1	8 .	80
/							

Table IX HLA-A3 Supermotif Peptides

HPV33	E1	9	80	HPV33	E2	11	284
HPV33		11	57	HPV33	E2	9	272
HPV33	E1	9	379	HPV33	E2	9	248
HPV33	E1	8	389	HPV33	E2	9	60
HPV33	E1	9	195	HPV33	E2	8	27
HPV33	E1	11	195	HPV33	E2	11	27
HPV33	E1	9	560	HPV33	E2	8	222
HPV33	E1	9	189	HPV33	E2	9	29
HPV33	E1	8	238	HPV33	E2	9	76
HPV33	E1	11	593	HPV33	E2	8	332
HPV33	E1	8	60	HPV33	E2	10	57
HPV33	E1	10	94	HPV33	E2	8	7
HPV33	E1	9	308	HPV33	E2	11	37
HPV33	E1	8	575	HPV33	E2	9	256
HPV33	E1	11	306	HPV33	E2	11	256
HPV33	E1	8	109	HPV33	E2	10	5
HPV33	E1	9	95	HPV33	E2	11	98
HPV33	E1	10	634	HPV33	E2	8	285
HPV33	E1	9	414	HPV33	E2	10	285
HPV33	E1	11	111	HPV33	E2	8	61 270
HPV33	E1	10	58	HPV33	E2	11	304
HPV33	E1	11	193	HPV33 HPV33	E2 E2	11 10	305
HPV33	E1	11	239	HPV33	E2 E2	11	209
HPV33	E1	8	447	HPV33	E2	11	254
HPV33	E1	9	447 558	HPV33	E2	8	257
HPV33	E1	11	328	HPV33	E2	10	257
HPV33 HPV33	E1 E1	10	240	HPV33	E2	8	310
HPV33	E1	8	299	HPV33	E2	10	233
HPV33	E1	8	491	HPV33	E2	8	118
HPV33	El	8	190	HPV33	E2	9	118
HPV33	E1	10	100	HPV33	E2	10	116
HPV33	E1	8	418	HPV33	E2	11	116
HPV33	E1	9	502	HPV33	E2	8	273
HPV33	El	9	522	HPV33	E2	9	117
HPV33	E1	9	595	HPV33	E2	10	117
HPV33	E1	11	254	HPV33	E2	9	58
HPV33	E2	8	249	HPV33	E2	11	58
HPV33	E2	9	258	HPV33	E2	10	102
HPV33	E2	9	245	HPV33	E2	11	102
HPV33	E2	8	40	HPV33	E2	9	309
HPV33	E2	10	288	HPV33	E2	11	159
HPV33	E2	9	211	HPV33	E5	9	12
HPV33	E2	10	25	HPV33	E5	11	10
HPV33	E2	8	235	HPV33	E5	11	38
HPV33	E2	9	143	HPV33	E5	9,	40
HPV33	E2	11	232	HPV33	E6	8	137
HPV33	E2	11	74	HPV33	E6		137 136
HPV33	E2	9	282	HPV33	E6	9	136
HPV33	E2	11	282	HPV33	E6	10	136
HPV33	E2	11	115	HPV33	E6		30
HPV33	E2	9	100	HPV33 HPV33	E6	10 11	98
HPV33	E2	10	156	HPV33	E6	8	27
HPV33	E2	10	278	HPV33	E6	9	27
HPV33	E2	9	15	HPV33	E6	9	47
HPV33	E2	11	4	HPV33	E6	11	45
HPV33	E2	10	14 165	HPV33	E6	9	69
HPV33	E2 E2	8	77	HPV33	E6	8	61
HPV33 HPV33	E2	9	284	HPV33	E6	10	99
HBA73	E2	,	204				

Table IX HLA-A3 Supermotif Peptides

HPV33 E	Ξ6	8	128	HPV33	L1	11	468
HPV33 E	36	9	64	HPV33	L1	11	62
HPV33 E	36	9	100	HPV33	L1	11	299
		8	70	HPV33	L1	9	57
HPV33 I	Ξ 6	10	25	HPV33	L1	10	221
HPV33 I	Ξ6	11	25	HPV33	L1	10	409
HPV33 E		9	127	HPV33	Ll	8	165
HPV33 E	Ξ6	8	86	HPV33	L1	11	55
HPV33 E		9	86	HPV33	L1	10	484
HPV33 E		8	109 -	HPV33	L1		484
		9	109	HPV33	L1	9	470
		8	95	HPV33	L1	11	470
		8	87	HPV33		8	156
		10	132	HPV33	L1	11	305
		10	68	HPV33	L1		254
		11	30	HPV33	L1	10	154
		8	59	HPV33	L1	11	328
		8	70	HPV33	L1		347
		10	31	HPV33 HPV33	L1 L1		481 488
		11	424	HPV33	L1	9	488
		8	411 44	HPV33	Li	11	488
		10 9	44 270	HPV33	L1		75
		9 11	207	HPV33	L1		455
		11	345	HPV33	L1		491
		8	103	HPV33	Li		491
		8	223	HPV33	L1		410
		9	457	HPV33	L1		51
		10	457	HPV33	L1	10	51
		10	370	HPV33	L1	10	32
		8	244	HPV33	L1	8	245
HPV33 I	L1	9	244	HPV33	L1		490
HPV33 I	-1	10	351	HPV33	L1		490
HPV33 I	L1	10	202	HPV33	Ll		227
HPV33 I		11	88	HPV33	L1		23
HPV33 I	51	10	269	HPV33	L1		486
		8	467	HPV33	L1		486
		10	467	HPV33	L1		486
		10	249	HPV33	L1		486 352
		10	50	HPV33 HPV33	L1 L1	9	301
		11	50	HPV33	L1	10	89
		8 11	256 419	HPV33	L1	11	31
		9	330	HPV33	L1	9.	421
		10	117	HPV33	Li	8	489
		9	472	HPV33	Li		489
		10	472	HPV33	Li		489
		8	68	HPV33	L1		485
		10	68	HPV33	L1	10	485
	51	11	226	HPV33	L1	11	485
HPV33 I	-1	9	118	HPV33	L1	11	10
HPV33 I	-1	10	425	HPV33	L1	10	66
HPV33 I	L1	8	474	HPV33	L1	9	22
HPV33 I	51	11	478	HPV33	L1	8	348
		8	429	HPA33	L1	10	300
		8	65	HPV33	L1		420
		11	65	HPV33	L1	8	331
		11	20	HPV33	L1	10	21
		11	43	HPV33 HPV33	L1 L1	10	101 12
HPV33 I	51	9	468	nrv33	nT.	9	12

Table IX HLA-A3 Supermotif Peptides

HPV33	L1	9	443		HPV45	E1	9	270
HPV33	L1	11	443		HPV45	E1	11	270
HPV33	L2	9	81		HPV45	E1	8	399
HPV33	L2	8	82		HPV45	E1	9	399
HPV33	L2	10	291		HPV45	E1	9	398
HPV33	L2	10	286		HPV45	E1	10	398
HPV33		11	286		HPV45	E1	8	297
HPV33		11	12		HPV45	E1	10	297
HPV33		9	308		HPV45	E1	11	423
HPV33		9	14		HPV45	E1	8	634
HPV33		8	447		HPV45	E1	9	78
HPV33	L2	9	447		HPV45	E1	10	78
HPV33		9	281		HPV45	E1	11	78
HPV33		11	301		HPV45	E1	10	214
HPV33		11	440		HPV45	E1	11	623
HPV33		11	58		HPV45	E1	8	328
HPV33	L2	11	226		HPV45	E1	8	596
HPV33		10	37		HPV45	E1	9	115
HPV33		10	25		HPV45	E1	10	115
HPV33		9	60		HPV45	E1	11	186
HPV33		10	379		HPV45	E1	8	189
HPV33		11	297		HPV45	E1	9	189
HPV33		11	285		HPV45	E1	10	295
HPV33		8	448		HPV45	E1	9	446
HPV33		9	292		HPV45	E1	8	456
HPV33		10	307		HPV45	E1	9	385
HPV33		11	311		HPV45	E1	10	449
HPV33		9	240		HPV45	E1	9	212
HPV33	L2	11	290		HPV45	E1	11	579
HPV33		10	215		HPV45	E1	8	19
HPV33	L2	8	444		HPV45	E1	9	19
HPV33		9	444		HPV45	E1	8	626
HPV33		10	444		HPV45	E1	9	209
HPV33		11	444		HPV45	E1	9	443
HPV33	L2	11	79		HPV45	E1	9	265
HPV33	L2	10	221		HPV45	E1	11	265
HPV33	L2	9	313		HPV45	E1	11	235
HPV33	L2	9	303		HPV45	E1	8	256
HPV33	L2	11	303		HPV45	E1	8	268
HPV33	L2	10	13		HPV45	E1	11	268
HPV33	L2	9	212		HPV45	E1	11	555
HPV33	L2	9	38		HPV45	E1	11	466
HPV33	L2	8	213		HPV45	E1	8	447
HPV33	L2	10	80		HPV45	E1	8	538
HPV33	L2	8	39		HPV45	E1	8	116
HPV33	L2	8	309		HPV45	E1	9	116
HPV33	L2	8	293		HPV45	E1	11	116
HPV33	L2	11	293		HPV45	E1	9	197
HPV33	L2	10	211		HPV45	E1	9	425
HPV33	L2	10	298		HPV45	E1	10	387
HPV33	L2	9	222		HPV45	E1	10	269
HPV33	L2	10	441		HPV45		8	299
HPV33	L2	11	441		HPV45	E1	9	267
HPV33	L2	11	210		HPV45	E1	11	84
HPV33	L2	9	228		HPV45	E1	8	190
HPV33	L2	8	381		HPV45	E1	8	271
HPV45	E1	11	383		HPV45	E1	10	271
HPV45	E1	8	198		HPV45	E1	10	556
HPV45	E1	11	532		HPV45	E1	10	467
HPV45	E1	11	452		HPV45	E1	8	210

Table IX HLA-A3 Supermotif Peptides

				-	•			
HPV45	E1	11	210		HPV45	E2	9	242
HPV45	E1	10	103		HPV45	E2	11	242
HPV45	E1	9	557		HPV45	E2	10	295
HPV45	E1	9	215		HPV45	E2	8	124
HPV45	E1	9	298		HPV45	E2	8	293
HPV45	E1	8	415		HPV45	E2	10	21
HPV45	E1	11	415		HPV45	E2	8	70
HPV45	E1	9	560		HPV45	E2	8	36
HPV45	E1	9	414		HPV45	E2	9	146
HPV45	E1	8	119		HPV45	E2	10	77
HPV45	E1	9	119		HPV45	E2	11	20
HPV45	El	10	119		HPV45	E2	9	232
HPV45	El	9	379		HPV45	E2	11	121
HPV45	E1	9	537		HPV45	E2	10	272
HPV45	E1	8	238		HPV45	E2	11	272
HPV45	E1	8	5 9 3		HPV45	E2	11	10
HPV45	E1	11	593		HPV45	E2	8	256
HPV45	E1	11	102		HPV45	E2	11	336
HPV45	E1	9	412		HPV45	E2	10	83
HPV45		11	412		HPV45	E2	8 .	46
HPV45	E1	8	80		HPV45	E2	9	69
HPV45		9	80		HPV45	E2	9	301
HPV45		8	451		HPV45	E2	11	301
HPV45		11	306		HPV45	E2	11	33
HPV45	E1	8	117		HPV45		9	109
HPV45		10	117		HPV45		10	109
	E1	11	117 307		HPV45 HPV45	E2 E2	9	289 292
HPV45		10 9	307		HPV45		8	67
HPV45 HPV45	E1	9	104		HPV45		11	67
HPV45		8	558		HPV45	E2	11	271
HPV45	E1	11	558			E2 -	10	112
HPV45	E1	11	239		HPV45	E2	9	35
HPV45		8	309		HPV45		11	222
HPV45		10	240		HPV45	E2	11	82
HPV45	E1	8	81		HPV45	E2	9	244
HPV45	E1	8	266		HPV45	E2	10	4
HPV45	E1	10	266		HPV45	E2	10	63
HPV45	E1	8	400		HPV45	E2	11	43
HPV45	E1	11	400		HPV45	E2	8	13
HPV45	E1	11	325		HPV45	E2	8	302
HPV45	E1	8	418		HPV45	E2	10	302
HPV45		9	502		HPV45	E2	11	302
HPV45	E1	9	522		HPV45	E2	8	275
HPV45	E1	10	254		HPV45		9	275
HPV45	E1	10	394		HPV45	E2	10	275
HPV45	E2	9	78		HPV45	E2	11	321 276
HPV45	E2	9	84		HPV45 HPV45	E2 E2	9	276
HPV45	E2 E2	8	305 274			E2	10	322
HPV45	E2 E2	9	274		HPV45	E2	9	51
HPV45		10	274		HPV45	E2	8	233
HPV45	E2	11	274		HPV45	E2	8	277
HPV45		11	158		HPV45	E2	8	290
HPV45	E2	9	171		HPV45	E2	11	290
HPV45		10	212		HPV45	E2	8	172
HPV45	E2	10	50 .		HPV45	E2	10	122
HPV45		9	255		HPV45	E2	9	213
HPV45	E2	8	225		HPV45	E2	10	337
HPV45	E2	8	242		HPV45	E2	8	214

Table IX HLA-A3 Supermotif Peptides

				-			
HPV45	E2	10	159	HPV45	L1	11	518
HPV45	E2	9	338	HPV45	L1	10	162
HPV45	E2	9	64	HPV45		8	164
HPV45		11	64	HPV45		11	88
HPV45		10	145	HPV45		10	276
HPV45		10	175	HPV45		8	129
HPV45		9	59	HPV45		11	188
HPV45		9	68	HPV45		8	250
HPV45		10	32	HPV45		9	488
HPV45		10	27		L1	10	488
HPV45		8	97	HPV45		9	271
HPV45		11	97 43	HPV45 I		11	271 114
HPV45		8 11	47	HPV45		10	296
HPV45		8	128	HPV45		11	169
HPV45		8	60	HPV45 I		8	283
HPV45		9	102	HPV45		8	24
HPV45		10	101	HPV45 I		8	498
HPV45		10	1			9	498
HPV45		8	100	HPV45 I		10	498
HPV45		11	100	HPV45 I	61	8	237
HPV45	E6	10 -	83	HPV45 I	L1	11	450
HPV45	E6	8	139	HPV45 I	L1	9	359
HPV45	E6	11	139	HPV45 I	L1	10	82
HPV45	E6	10	95	HPV45 I	61	11	503
HPV45	E6	9	114	HPV45 I		10	143
HPV45		11	114	HPV45 I		11	328
HPV45		8	111	HPV45 I		10	473
HPV45		9	111	HPV45 I		8	91
HPV45		8	144	HPV45		11	91
HPV45		9	144	HPV45 I		10	68
HPV45		10	144	HPV45 I		9 9	144 69
HPV45		11	144	HPV45 I		8	499
HPV45		10	41 29	HPV45 I		9	499
HPV45		9	84	HPV45 I		8	49
HPV45		9	28	HPV45 I		9	383
HPV45		8	72	HPV45 I			516
HPV45		9	64	HPV45 I		9	190
HPV45		8	78	HPV45 I		8	526
HPV45		10	44	HPV45 I	51	10	526
HPV45		11	44	HPV45 I	51	11	526
HPV45	E7	8	47	HPV45 I	51	10	22
HPV45	E7	11	62	HPV45 I	51	10	248
HPV45	E7	8	61	HPV45 I			508
HPV45	E7	11	75	HPV45 I			387
HPV45		9	51	HPV45 I			440
HPV45		11	51	HPV45 I			380
HPV45		11	49	HPV45 I			281
HPV45		8	54	HPV45 I			334
	E7	10	76	HPV45 I			357
HPV45		9	45	HPV45 I		9	452 67
HPV45		10	45	HPV45 I			101
HPV45 HPV45		9 11	517 161	HPV45 I			529
HPV45		8	191	HPV45 I		11	46
HPV45		11	234	HPV45 I			77
HPV45		9	523	HPV45 I			93
HPV45		11	523	HPV45 I			58
HPV45		8	518	HPV45 I			272
	-						

Table IX HLA-A3 Supermotif Peptides

HPV45	L1	11	486	HPV45	L2	10	302
HPV45	L1	8	521	HPV45	L2	11	298
HPV45	L1	9	521	HPV45	L2	10	281
HPV45	L1	11	521	HPV45	L2	11	281
HPV45	L1	10	115	HPV45	L2	11	225
HPV45	L1	10	519	HPV45		9	308
HPV45	L1	11	519	HPV45	L2	10	68
HPV45	Ll	8	453	HPV45	L2	10	220
HPV45	L1	8	522	HPV45	L2	10	13
HPV45	L1	10	522	HPV45	L2	8	288
HPV45		9	163	HPV45	L2	11	288
HPV45	L1	9	116	HPV45	L2	9	211
HPV45	L1	9	330	HPV45	L2	10	211
HPV45	L1	11	57	HPV45	L2	11	362
	L1	8	442	HPV45	L2	8	212
HPV45		9	520	HPV45	L2	9	212
HPV45	L1	10	520	HPV45	L2	11 9	358
	L1	10	329	HPV45	L2 L2	10	363
HPV45	L1	9	441	HPV45	L2	8	304
	L1	8	70	HPV45	L2	8	359
HPV45		9	297	HPV45	L2	10	293
	L1	11	36 102	HPV45	L2	9	217
HPV45 HPV45	L1 L1	10	92	HPV45	L2	9	80
HPV45		8	360	HPV45	L2	8	2
HPV45	L1	10	47	HPV45	L2	9	2
HPV45	L1	9	78	HPV45	L2	10	2
	Li	10	127	HPV45	L2	8	81
	L1	8	196	HPV45	L2	11	437
	L1	8 .	477	HPV45	L2	9	227
HPV45	L1	10	303	HPV56	E2	9	177
HPV45	Li	9	38	HPV56	E2	10	177
HPV45	L2	10	286	HPV56	E2	8	178
	L2	8	12 :	HPV56	E2	9	178
HPV45		11	12	HPV56	E2	11	178
HPV45	L2	10	357	HPV56	E2	8	4
HPV45	L2	9	14	HPV56	E2	8	71
HPV45	L2	9	303	HPV56	E2	10	176
HPV45	L2	9	273	HPV56	E2	11	176
HPV45	L2	9	276	HPV56	E2	9	195
HPV45	L2	11	306	HPV56	E2	8	140
	L2	11	58	HPV56	E2	8	213
HPV45	L2	10	25	HPV56	E2	10	213
	L2	9	60	HPV56	E2	8	117 43
HPV45	L2	11	292	HPV56 HPV56	E2 E2	9	43
	L2	10	210	HPV56	E2	8	191
HPV45	L2	11	210	HPV56	E2	10	154
HPV45	L2	10	34	HPV56	E2	8	61
HPV45	L2	9	287	HPV56	E2	10	99
HPV45	L2	9	1	HPV56	E2	10	59
HPV45	L2 L2	10	1	HPV56	E2	11	210
		11	1	HPV56	E2	8	239
HPV45 HPV45	L2 L2	10	79	HPV56	E2	10	239
HPV45	L2 L2	11	285	HPV56	E2	9	297
HPV45	L2	11	356	HPV56	E2	9	283
HPV45	L2	11	209	HPV56	E2	10	211
HPV45	L2	9	214	HPV56	E2	11	281
HPV45	L2	10	216	HPV56	E2	9	233
	L2	9	11	HPV56	E2	9	90
-11 4-4-3		-	-				

Table IX HLA-A3 Supermotif Peptides

					-			
HPV56	E2	11	295		HPV56	E6	10	135
HPV56	E2	9	46		HPV56	E6	8 .	73
HPV56	E2	10	46		HPV56	E7	10	75
HPV56	E2	9	1		HPV56	E7	8	39
HPV56	E2	11	1		HPV56	E7	11	70
HPV56	E2	8	292		HPV56	E7	10	42
HPV56	E2	11	236		HPV56	E7	8	62
HPV56	E2	10	301		HPV56	E7	10	60
HPV56	E2	8	246		HPV56	E7	8	73
HPV56	E2	11	246		HPV56	E7	8	77
HPV56	E2	11	188		HPV56	E7	10	71
HPV56	E2	8	279		HPV56	E7	11	59
HPV56	E2	11	223		HPV56	L1	11	241
HPV56	E2	8	196		HPV56	L1	8	198
HPV56	E2	11	266		HPV56	L1	8	58
	E2	10	282		HPV56	L1	11	381
HPV56	E2	11	28		HPV56	L1	9	444
HPV56	E2	8	234		HPV56	L1	9	37
HPV56	E2	9	155		HPV56	L1	8	512
HPV56	E2	8	179		HPV56	L1	11	79
HPV56	E2	10	179		HPV56	L1	11	195 136
HPV56	E2	10	237		HPV56 HPV56	L1 L1	10 '	389
HPV56	E2	9	302		HPV56	LI	8	257
HPV56	E2	10	45 45		HPV56	L1	9	491
HPV56	E2	11	278		HPV56	L1	10	491
HPV56	E2 E2	9	111		HPV56	L1	8	278
HPV56	E2	10	111		HPV56	L1	9	278
HPV56	E6	8	89	-	HPV56	L1	11	176
HPV56	E6	9	89		HPV56	Li	10	236
HPV56	E6	10	139		HPV56	Li	11	121
HPV56	E6	9	69		HPV56	L1	10	404
HPV56	E6	10	69		HPV56	L1	10	308
HPV56	E6	9	50		HPV56	L1	10	303
HPV56	E6	10	33		HPV56	L1	8	290
HPV56	E6	8	101		HPV56	L1	11	290
HPV56	E6	10	28		HPV56	L1	8	501
HPV56	E6	11	28		HPV56	L1	10	501
HPV56	E6	8	23		HPV56	L1	8	33
HPV56	E6	11	20		HPV56	L1	9	364
HPV56	E6	11	44		HPV56	L1	10	150
HPV56	E6	11	48		HPV56	L1	10	378
HPV56	E6	9	88		HPV56	L1	10	334
HPV56	E6	10	88		HPV56	L1	8	98
HPV56	E6	8	137		HPV56	L1	11	98
HPV56	E6	8	70		HPV56	L1	10	55
HPV56	E6	9	70		HPV56	L1	11	55
HPV56	E6	11	70		HPV56	L1	11	45
HPV56	E6	8	31		HPV56	L1	11	168
HPV56	E6	8	98		HPV56	Ll	9	502 151
HPV56	E6	11	98		HPV56	L1	9	385
HPV56	E6	8	119		HPV56 HPV56	L1 L1	10	365
HPV56	E6	9	119		HPV56	PT PT	11	333
HPV56	E6	9	110		HPV56	L1	9	333
HPV56	E6	8	30		HPV56	L1	10	1
HPV56	E6	9	30		HPV56	L1	11	95
HPV56 HPV56	E6 E6	9 11	67 67		HPV56	L1	9	123
HPV56 HPV56		8	90		HPV56	Ll	8	91
HPV56	E6	10	21		HPV56		9	197
nrvob	50	10					-	

Table IX HLA-A3 Supermotif Peptides

HPV56	L1	8	511	HPV56 L1		134
HPV56	Ll	9	511	HPV56 L1		203
HPV56		10	31	HPV56 L1		310
HPV56		10	255	HPV56 L1		47
HPV56		10	467	HPV56 L1		283
HPV56		9 .	522	HPV56 L1		85
HPV56		10	522	HPV56 L1		453
HPV56		11	522	HPV56 L2		222
HPV56		9	442	HPV56 L2		281
HPV56		11	442	HPV56 L2		281
HPV56		10	288	HPV56 L2		438
HPV56		11	339	HPV56 L2		438
HPV56		11	362	HPV56 L2		438
HPV56		8	384	HPV56 L2		12
HPV56		9	384	HPV56 L2		12
HPV56		11	35	HPV56 L2 HPV56 L2		367 14
HPV56		11	260	HPV56 L2		30
HPV56		10	508	HPV56 L2		437
HPV56		11	508			437
HPV56		9	455	HPV56 L2 HPV56 L2		276
HPV56		9 '	108	HPV56 L2		287
HPV56		11	520	HPV56 L2		58
HPV56		9	100	HPV56 L2		25
HPV56		11	100			60
HPV56		10	67	HPV56 L2 HPV56 L2		293
HPV56		10	446	HPV56 L2		293
HPV56		8	279	HPV56 L2		221
HPV56			456	HPV56 L2		221
HPV56		9	379	HPV56 L2		210
HPV56		10	261 489	HPV56 L2		210
HPV56		11	526	HPV56 L2		81
HPV56		9	526	HPV56 L2		302
			524	HPV56 L2		34
HPV56 HPV56		9	524	HPV56 L2		235
HPV56		10	524	HPV56 L2		1
HPV56		11	524	HPV56 L2		1
HPV56		8	86	HPV56 L2		î
HPV56		8	380	HPV56 L2		ī
HPV56		9	304	HPV56 L2		285
HPV56		11	377	HPV56 L2		209
HPV56		9	335	HPV56 L2		369
HPV56		11	66	HPV56 L2	11	78
HPV56		8	445	HPV56 L2	8	441
HPV56		11	445	HPV56 L2	9	441
HPV56		8	525	HPV56 L2	10	441
HPV56		9	525	HPV56 L2	11	441
HPV56		10	525	HPV56 L2	11	306
HPV56		8	523	HPV56 L2	10	68
HPV56		9	523	HPV56 L2	9	11
HPV56		10	523	HPV56 L2	9	220
HPV56		11	523	HPV56 L2	10	220
HPV56		8	57	HPV56 L2	11	298
HPV56		9	57	HPV56 L2	11	225
HPV56		8	443	HPV56 L2	10	13
HPV56		10	443	HPV56 L2	9	211
HPV56		10	99	HPV56 L2		211
HPV56		8	365	HPV56 L2	10	79
HPV56		9 -	56	HPV56 L2		212
HPV56		10	56	HPV56 L2	9	212

Table IX HLA-A3 Supermotif Peptides

HPV56	L2	9	80
HPV56	L2	8	288
HPV56	L2	11	288
HPV56	L2	8	2
HPV56	L2	9	2
. HPV56	L2	10	2
HPV56	L2	11	280
HPV56	L2	11	366
HPV56	L2	9	236
HPV56	L2	8	31

SF 1168096 v1

2	3	4		L2	8	274
Ll	8 .	234		L2	11	274
E1	8	206		E1	10	143
L1	9	489		E1	8	336
Ll	11	489		E1	8	180
L2	10	286		E1	8	62
E1	9	112		E1	10	100
E1	10	112		E1	9	375
Ll	11	420		E1	10	105
E1	10	475		E6	8	42
L1	11	203		E6	11	42
L1	8	487		L1	9	453
L1	9	487		L1	10	453
L1	11	487		E1	9	197
L2	11	12		E1	8	604
E2	8	322		E2	11	74
L2	8	288		E1	8	417
L2	11	288		E2	8	100
L1	8	22		E2	9	100
E1	8	407		E1	11	373
L2	9	14		E2	9	293
E6	9	10		E2 E2	11 9	293 39
E6	9	86		E7	11	39
E1	8	77		E6	11	113
E1	9	77		L1	8	206
E1	9	101		LI	8	252
L1	10	43		L2	8	442
L1	11	43 231		E1	9	220
E2	8			E6	9	126
E2 E2	9 11	231 231		E6	11	126
L1	10	483		E1	9	454
Li	11	483		Li	8	463
E1	11	601		L1	10	463
E6	10	64		E4	11	21
L1	11	157		E1	9	393
E1	9	406		L1	10	245
E2	8	296		L2	9	276
E2	11	35		L1	8.	49
L1	8	99		E1	11	587
E1	8	640		Ll	9	326
E1	10	111		L2	11	58
E1	11	111		E1	9	194
E2	8	230		E2	10	156
E2	9	230		E1	8	350
E2	10	230		E1	9	217
L1	8	219		L2	11	292
E6	8	96		E2	9	55
E1	9	570		E1	9	273
E7	10	88		E1	11	273
E1	10	222		L1	11	130
E2	10	25		E1	11	431
E1	9	203		L2	9	303
E1	11	203		E1	8	632
L1	11	84		E1	10	191
E1	11	73		L2	10	25
L1	11	269		L2	9	60
E6	11	99		E1	8	145
E1	10	178		L1	8	407
E2	9	174		E4	9	90

E1	9	316		E1	11	540
Ll	9	478		E4	10	8
L1	10	113		E4	11	8
E1	10	415		E4	8	24
E2	11	53		E1	8	198
E6	10	119		L1	9	464
E4	8	10		E1	8	276
E4	9	10		E1	11	276
E1	9	246		E1'	11	563
L2	8	3		E1	8	218
L2	9	3		E1	11	218
E6	11	25		L1	9	439
L2	11	306		L1	11	439
L2	8	149		E4	8	81
E2	9	29		E6	8	121
E1	8	376		E1	9	115
E1	11	474		E1	10	115
L2	9	287		E1	11	115
L1	8	272		E1	10	277
E1	8	195		E1 :	8	279
E1	11	195		E1	10	279
E6	9	120		L2	10	293
E1	9	106		L1	11	295
E2	11	315		E1	10	564
L1	10	421		E7	11	67
E1	8	571		L1	9	233
E2	10	267		L2	8	1
E2	11	267		L2	9	1
L2	8	82		L2	10	1
E7	9	89		L2	11	1
E1	9	476		E1	8	546
L1	8	486		E1	8	421
L1	9	486		E1	10	421
L1	10	486		E1	8	274
E1	9	433		E1	10	274
E2	9	351		E1	10	607
E6	9	128		E1	9	568 568
E1	8	114		E1 E1	11 10	451
E1	10	114		L1	10	31
E1	11	114		E4	9	23
E1	9	462		L1	10	438
E1	10 9	462 420		E1	8	397
E1 E1	11	420		L1	9	352
E1	8	286		E1	11	59
E2	9	165		E1	10	395
E2	9	147		L2	9	38
E6	8	116		E2	10	348
E6	10	116		L2	9	237
E1	10	121		L2	11	285
E1	11	283		L1	11	482
Li	8	61		L2	8	438
L1	11	61		L2	9	438
L1	11	19		L2	10	438
Ll	9	71		Li2	11	438
L1	11	42		L1	10	217
Ll	9	271		E6	9	110
E6	9	101		E4	8	34
E1	9	223		E4	9	34
E2	11	266		L1	8	160

L1	10	160	L1	9	348
E2	8	61	E2	8	40
E1	9	545	E6	10	40
L2	10	80	E1	9	192
L1	9	266	E1	11	192
L2	8	212	L2	. 8	39
L2	10	212	L1	10	270
E4	9	38	E6	10	26
E4	10	3.8	E6	11	8
E4	11	38	L2	10	147
Еб	8	28	E1	8	566
L1	11	301	E1	11	566
L1	11	324	· E4	11	78
L1	10	232	L1	11	346
L1	10	250	E2	11	97
E2	9	76	E6	11	39
E1	8	305	E6	8	11
E1	11	314	E6	8	87
L1	11	466	E4	8	91
L1	9	417	E1	8	78
E2	9	103	E2	10	334
E2	10	103	E1	8	317
E2	11	103	L1	9	21
E2	9	233	E1	11	333
E1	9	205	E2	8	30
E1	11	391	E6	10	100
L1	9	53	E4	11	7
L2	11	298	E1	9	275
E1	9	109	E1	9	278
L1	8	241	E1	11	278
L2	10	281	L1	9	114
L2	11	281	L1	10	62
L2	9	308	L1	9	484
L1	9	472	L1	10	484
L1	10	472	L1	11	484
L1	11	476	L1	10	296
L1	9	140	L2	11	146
L1	8	488	E1	9	565
L1	10	488	E2	11	333
L2	10	13	L1	8	327
E6	10	9	E2	9	335
E1	9	422	L1	10	20
L1	8	474	E2	9	349
E1	8	247	E2	11	349
L2	9	81	Ll	8	72
E1	10	60	L1	11	58
L1	9	86	E2	11	58
E4	9	80	E7	10	68
L2	8	304	L1	10	97
L1	9	297	E2	9	321
L1	11	451	E1	8	426
L1	8	473	E1	9	530
L1	9	473	E1	8	464
L1	10	85	El	9	510
E4	10	79	E2	11	145
L2	11	209	E6	10	85
E2	10	316	E1	8	76
L1	10	347	E1	9	76
L2	10.	210	E1	10	76
E2	9	317	E6	11	46

L2	10	435
L2	11	435
E6	9	44
L1	11	350
E1	10	402
E2	10	168
L2	9	71
L1	10	10
E2	10	138
L1	11	415

Table IXB. HPV6B HLA-A3 Supermotif Peptides

				•	•	
2	3	4			2 9	174
L1	8	234		I	.2 8	274
E1	8	206		I	.2 1	1 274
Ll	9	489		E	1 1	.0 143
L1	11	489		E	1 8	336
L2	10	286		E	1 8	180
El	9	112		E	1 8	62
E1	10	112		E	1 1	0 100
L1	11	420		E	1 9	375
E1	10	475		E	1 1	0 105
L1	11	203		E	6 8	42
L1	8	487		E	6 1	1 42
L1	9	487		I	1 9	453
L1	11	487		L	1 1	0 453
L2	11	12		E	1 9	197
L2	8	288		E	1 8	604
L2	11	288		E	2 1	1 74
E2	8	322		E	1 8	417
L1	8	22		E	2 8	100
E1	8	407		E	2 9	100
L2	9	14		E	1 1	1 373
E6	9	10		E	2 9	293
E6	9	86		E	2 1	1 293
E1	8	77		E	2 9	39
E1	9	77		E	7 1	1 39
E1	9	101		E	6 1	1 113
L1	10	43		L	1 8	206
L1	11	43		L	1 8	252
E5B	9	36	-	L	2 8	442
E5B	11	36		E	1 9	220
E2	8	231		E	6 9	
E2	9	231		E	6 1	1 126
E2	11	231			1 9	
L1	10	483			1 8	
Ll	11	483				0 463
E1	11	601		E		
E6	10	64			1 9	
L1	11	157		L		
E1	9	406		L		276
E2	8	296		L		
E2	11	35		E		
L1	8	99		L		326
E1	8	640		L		
El	10	111		E		194
E1	11	111		E		
E2	8	230		E		350
E2	9	230			5B 8	28
E2	10	230		E		
L1	8	219		L		
E6	8	96			5B 1	
E1	9	570		E		55
E2	10	25		E		
E1	10	222		E		
El	9	203		L		
E1	11	203		E		
Ll	11	84		L		303
E1	11	73		E		632
L1	11	269		Е		
E6	11	99		L		
E1	10	178		L	2 9	60

Table IXB. HPV6B HLA-A3 Supermotif Peptides

			HLA-A3 Supermotif	Peptides		
E1	8	145		L1	9	271
L1	8	407		E6	9	101
E4	9	100		E1	9	223
E1	9	316		E2	11	266
Ll	9	478		E1	11	540
Li	10	113		E4	10	18
E1	1:0	415		E4	11	18
E2	11	53		E4	8	34
E6	10	119		E1	8	198
E4	8	20		L1	9	464
E4	9	20		E1	8	279
E1	9	246		E1	10	279
L2	8	3		E1	8	276
L2	9	3		E1	11	276
E5B	9	42		E1	11	563
E6	11	25		E5B	11	31
L2	11	306		E1	8	218
L2	8	149		E1	11	218
E2	9	29		L1	9	439
E1	8	376		L1	11	439
E1	11	474		E4	8	91
L2	9	287		E6	8	121
L1	8	272		E1	9	115
E1	8	195		E1	10	115
E1	11	195		E1	11	115
E6	9	120		E1	10	277
E1	9	106		L2	10	293
E2	11	315		L1	11	295
L1	10	421		E1	10	564
E1	8	571		E7	11	67
E2	10	267		L1	9	233
E2	11	267		L2	8	1
L2	8	82		L2	9	1
E7	9	89		L2	10	1
E1	9	476		L2	11	1
L1	8	486		E5B	10	26
L1	9	486		E1	8	546
L1	10	486		E1	8	421
E1	9	433		E1 E1	10	421 274
E2	9	351		E1	10	274
E6	9	128		E1	9	568
E1	8	114		E1	11	568
E1	10	114		E1	10	451
E1	11	114		L1	10	31
E1 E1	9 10	462 462		E7	10	88
E1		420		E4	9	33
E1	9 11	420		L1	10	438
E1	8	286		E1	8	397
E2	9	165		Ll	9	352
E2	9	147		E1	11	59
E6	8	116		E1	10	395
E6	10	116		L2	9	38
E1	10	121		E1	10	607
E1	11	283		L2	9	237
LI	8	61		L2	11	285
L1	11	61		Li	11	482
Li	11	19		L2	8	438
L1	9	71		L2	9	438
Li	11	42		L2	10	438
-						

Table IXB. HPV6B HLA-A3 Supermotif Peptides

			rica-A3 Supermoni	repudes		
L2	11	438		L1	9	473
L1	10	217		L1	10	85
E6	9	110		E4	10	89
E4	8	44		L2	11	209
E4	9	44		E2	10	316
L1	8	160		L1	10	347
L1	10	160		E2	9	220
E2	8	61		L2	10	210
E1	9	545		E2	9	317
L2	10	80		L1	9	348
L1	9	266		E2	8	40
L2	8	212		E6	10	40
L2	10	212		E1	9	192
E4	9	48		E1	11	192
E4	10	48		L2	8	39
E4	11	48		L1	10	270
E6	8	28		E6	10	26
L1	11	301		E2	8	221
L1	11	324		E6	11	8
L1	10	232		L2	10	147
L1	10	250		E1	8	566
E2	9	76		E1	11	566
E1	8	305		E4	11	88
E1	11	314		L1	11	346
L1	11	466		E2	10	219
L1	9	417		E2	11	97
E2	9	103		E6	11	39
E2	10	103		E6	8	11
E2	11	103		E6	8	87
E2	9	233		E4	8	101
E2	11	218		E1	8	78
E1	9	205		E2	10	334
E1	11	391		E1	8	317
L1	9	53		Ll	9	21
E5B	10	35		E1	11	333
L2	11	298		E2	8	30
E1	9	109		E6	10	100
L1	8	241		E4	11	17
L2	10	281		E1		278
L2	11	281		E1		278 275
L2	9	308		E1		114
L1	9	472		L1 L1	9 10	62
L1	10	472				484
L1	11	476				484
L1	9	140 488				484
L1	8 10	488				296
L1 L2	10	13		L2		146
E6	10	9				565
E1	9	422				333
L1	8	474				327
B1	8	247		E2		335
L2	9	81		L1		20
E1	10	60				349
L1	9	86				72
E4	9	90				58
L2	8	304				58
L1	9	297				68
L1	11	451				97
L1	8	473		E2	9	321
	-					

Table IXB. HPV6B HLA-A3 Supermotif Peptides

E1	8	426
E1	9	530
E1	8	464
E1	9	510
E2	11	145
E6	10	85
E1	8	76
E1	9	76
E1	10	76
E6	11	46
L2	10	435
L2	11	435
E6	9	44
L1	11	350
E1	10	402
E2	10	168
L2	9	71
L1	10	10
E2	10	138
L1	11	415

Table IXC HPV11 HLA-A3 Supermotif Peptides

				-	-		
2	3	4			E1	8	336
L1	8	235			E1	8	62
E1	9	112			E1	8	180
E1	10	112			E1	9	375
E1	8	407			E6	11	113
L1	11	421			E1	10	105
L2	9	80			E6	8	42
L2	10	285			E6	11	42
E1	10	475			L1	9	454
L1	11	204			L1	10	454
L2	11	11			E1	9	197
E2	8	321			E6	10	69
E2	8	197			E1	8	604
L1	8	22			E1	8	417
L2	9	13			E2	11	74
E6	9	10			E2	8	100
E1	8	77			E2	9	100
E1	9	77			E1	11	373
L1	9	349			E2	9	39
E1	9	101			E6	8	92
L1	10	43			E1	10	141
Ll	11	43			L1	8	207
E5	9	37			L1	8	253
E5	11	26			L2	8	438
Ll	10	484			E6	9	126
L1	11	484			E6	11	126
E1	11	601			E1	9	454
E6	10	64			L1	8	464
E1	9	406			L1	10	464
E5	8	46 '			E1	9	393
L1	11	158			L2	9 11	275 587
E2	11	35			E1	9	220
E2	8	295			E1	9	327
E2	11	194			L1	11	57
L1	8	99			L2 L1	9	479
E1	8	640			E1	9	194
E1	11	73 111			E2	10	156
E1	10	111			E5	8	29
E1 E1	11 10	607			E1	9	217
L1	8	220			L2	11	291
E1	10	168			E1	9	273
E2	10	25			E1	11	273
E6	8	96			E1	8	632
E1	9	203			L2	10	24
E1	11	203			E5	9	40
E1	9	570			E5	11	40
E1	10	222			L2	9	59
L1	10	271			E1	11	431
L1	10	439			Ll	8	408
E1	9	46			E4	9	99
E2	9	292			E1	9	316
E2	11	292			E2	8	232
E7	9	31			E2	9	232
L1	11	84			E2	11	232
E1	10	191			L1	10	113
E1	8	143			E1	10	415
E1	10	178			E6	10	119
L2	8	273			E2	9	29
L2	11	273			L1	11	325

Table IXC HPV11 HLA-A3 Supermotif Peptides

PCT/US00/33549

			HLA-A3 Supermotif	Peptides		
E4	9	20		L1	9	465
E1	8	305		Ll	9	272
E1	9	246		E1	8	276
E1	9	349		E1	11	276
E6	11	25		E1	11	563
L2	10	36		E1	8	218
E1	8	376		E1	11	218
E1	11	474		E4	8	90
E1	8	195		E2	9	103
E1	11	195		E2	10	103
E6	9	120		E1	9	115
E2	8	30		E1	10	115
E1	8	571		E1	11	115
E1	9	106		E2	9	334
L1	10	422		L1	8	273
L2	8	81		E1	10	277
L2 L2	9	286		L2	10	292
	9	89		L1	11	296
E7	9	476		E1	10	564
E1	8			E7	11	67
E2		265				
E2	11	265		L1 L1	9	234
E2	9	350		L1	11	440 440
E6	8	116		E1	8	546
E6		116				
E1	11	460		E1	8	421
E6	9	128		E1	10	421
E1	8	114		L1	9	339
E1	10	114		E1 E1	8	47 274
E1	11	114		E1		274
E1.	9	462		E4	10	1
E1	10	462		E4 E1	10	100
E1	9	420			9	568
E1	11	420 347		E1 E1	11	568
L1	11			E1	10	451
E1	10	484		L1	10	31
E1 E5	8	286 43		E7	10	88
	8 11	43		E2		264
E5 E2	9	165		E4		33
	9	433		E1	8	397
E1 E1	10	121		L1		338
E6	11	99		L1	9	353
E1	11	283		E1	10	395
L1	9	53		E1		59
	8	61		E2		347
L1 L1	11	61		E2		248
E2	9	147		L2		236
L1	10	19				79
L1	11	19		L2		284
Ll	9	71				49
L1	11	42				49
E6	9	101		E4		49
E1	8	279				196
E1	10	279				483
E1	9	279				110
		540				218
E1 E4	11	18				161
						211
E4	8	34 198				211
E1 E4	11	31				44
£4	11	2.7	1		-	**

Table IXC HPV11 HLA-A3 Supermotif Peptides

				•		
E4	10	44		E2	11	348
E2	8	61		E2	8	40
E1	9	545		E6	10	40
L1	8	488		L1	9	490
L1	9	488		L1	11	490
L1	11	488 .		L2	8	38
L2	9	206		L1	8	340
L2	8	434		E2	8	249
L2	9	434		E6	10	26
L2	10	434		E2 L1	10 8	333 474
L2	11	434		LI	9	474
L1	9	267 28		E6	11	8
E6 L1	11	302		E1	9	192
E2	10	288		E1	11	192
L1	10	233		E6	8	11
L1	10	251		E1	8	566
E2	9	76		E1	11	566
El	11	314		L2	8	287
L1	11	467		L2	11	287
L1	9	418		E4	11	87
E1	11	391		L2	8	207
E5	8	36		E2	11	97
E5	10	36		E6	11	39
L2	11	297			11	228
E2	9	37			9	37
E2	11	37		E1	8	116
L1	8	242		E1	9	116
E4	9	59			10	116
L2	10	280		E4	8	100
L2	11	280			8	78 21
L2	9	307		L1	8 9	21
E1	9	205 477		E1	11	333
L1 L1	11 9	473		E5	10	27
L1	10	473		E6	10	100
LI	9	141		E1	9	278
E2	9	234			11	278
L1	8	475			9	275
L2	10	12		E2	8	233
E6	10	9		E2	10	233
L1	10	348			9	114
E1	8	247			10	62
E1	9	422			9	485
E1	8	350				485
L1	9	86				485
E4	9	89				297
L1	9	298				266
E7	8	32				266
E7	11	32			11 9	332 565
L1	11	452			8	148
L1	10	85 88			8	328
E4 L2	10	208			9	20
L1	8	489			10	20
L1	10	489		E7	10	68
E1	8	206			8	72
L2	10	209			11	58
El	10	60			11	58
E2	9	348		L1	10	97
-						

Table IXC HPV11 HLA-A3 Supermotif Peptides

E2	9	320
E1	8	426
E1	8	464
E1	9	510
E2	10	102
E2	11	102
E1	. 9	530
E2	11	145
E1	8	76
E1	9	76
E1	10	76
E6	9	44
E6	11	46
L2	9	70
L2	10	431
L2	11	431
E2	10	138
L1	10	246
L1	8	49
L1	11	351
L2	11	305
E1	10	402
E2	10	168
L1	10	10
L1	11	416

Table X HLA A24 Supermotif Peptides

				HLA AZ4	supermour	replices	•		
1 :	2	3	4			HPV16	E1	10	392
		8	240			HPV16	E1	11	463
HPV16	E1	11	391		1	HPV16	E1	9	493
	E1	9	539		1	HPV16	E1	9	445
HPV16	E1	9	459		1	HPV16	E1	10	445
HPV16	E1	9	318		1	HPV16	E1	8	456
HPV16	E1	9	206			HPV16	E1	9	456
HPV16	E1	10	206		1	HPV16	E1	8	453
HPV16 F	E1	8	524		3	HPV16	E1	11	453
HPV16	E1	9	524		1	HPV16	E1	9	585
HPV16	E1	9	82			HPV16	E1	8	501
HPV16 H	E1	10	82			HPV16	E1	9	501
HPV16	E1	11	23			HPV16	E1	10	501
		9	500			HPV16	E1	11	508
HPV16 E	E1	10	500			HPV16	E1	8	466
		11	500			HPV16	E1	9	466
		11	237			HPV16	E1	10	466
		9	259			HPV16	E1	11	466
HPV16 F		9	304			HPV16	E1	9	325
		8	353			HPV16	E1	10	242
		11	353			HPV16	E1	11	519
HPV16 E	31	10	101			HPV16	E1	8	487
HPV16 E		10	640			HPV16	E1	10	272
HPV16 H		8	299			HPV16	E1	8	571
		9	299			HPV16	E1	9	571
		9	528			HPV16	E1	8	12
		9	50			IPV16	E1	9	12
		10	50			IPV16	E1	8	450
		10	97			HPV16	E1	11	450
		8	368			IPV16	E1	8	179
		10	368			IPV16	E1		216
		9 .	43			IPV16	E1		263
HPV16 E		10	43			HPV16	E1		263
		9	384			HPV16	E1 E1		184
		10	384			HPV16	E1	11	411
		10	548			HPV16		9	369
			235			HPV16	E1 E1	11	369
HPV16 F		8	438			HPV16	E1	8	401
		9	438			HPV16	E1		52
		11 9	438 452			IPV16	E1		210
		9	374			HPV16	E1	8	492
			603			IPV16	E1	10	492
		10 11	603			IPV16	E1	9.	400
			356			IPV16	E1		296
			356			IPV16	E1		296
			213			IPV16	E1		296
		8	63			IPV16	E1		311
		9	63			IPV16	El		311
			152			IPV16	El	11	311
			288			IPV16	E1	11	323
			288			IPV16	El		252
		8	138			IPV16	E1		252
		9	331			IPV16	E1		252
			331			IPV16	E1	9	199
			338			IPV16	E1	10	199
		9	612			IPV16	E1	10	89
		11	612			IPV16	E1	9	126
			51			IPV16	E1	9	485
		9	51			IPV16	E1	10	485
E A TO E		-							

Table X HLA A24 Supermotif Peptides

				TEST TEST Supermons	repude			
HPV16	E1	9	297		HPV16	E1	8	343
HPV16	E1	10	297		HPV16	E1	8	84
HPV16		11	297		HPV16	E1 '	8	362
HPV16	E1	8	254		HPV16	E1	8	257
HPV16		9	254		HPV16	E1	11	257
HPV16		11	254		HPV16		10	125
HPV16		11	293		HPV16		11	582
HPV16		10	490		HPV16		8	615
HPV16		8	457		HPV16	E1	8	432
HPV16		11	457			E1	11	432
HPV16		8	191		HPV16		9	575
HPV16		9	243		HPV16	E1	11	280
HPV16		11	48		HPV16		8	447
HPV16		9	554		HPV16		10	447
HPV16		10	554		HPV16		11	447
HPV16		11	554		HPV16		10	611
HPV16		11	544		HPV16	E1	9	455
HPV16		8	91		HPV16		10	455
HPV16		10	583		HPV16		9	349
HPV16		11	475		HPV16		9	218
HPV16		9	214		HPV16		9	246
HPV16		8	260		HPV16		10	246
HPV16		8	319		HPV16		9	250
HPV16		11	319		HPV16		11	250
HPV16		10	444		HPV16		8	266
HPV16		11	444		HPV16		11	266
HPV16		8	207		HPV16		8	484
HPV16		9	207		HPV16		10	484
HPV16		10	520		HPV16		11	484
HPV16		8	305		HPV16		11	489
HPV16		10	360		HPV16	E1	9	546
HPV16		9	273		HPV16	E1	8	421
HPV16		10	567		HPV16	E1	8	314
HPV16		9	105		HPV16	E1	9	314
HPV16		11	105		HPV16	E1	8	231
HPV16		10	535		HPV16	E1	11	231
HPV16		9	136		HPV16	E1	8	270
HPV16		10	136		HPV16	E1	9	270
HPV16		9	480		HPV16	E1	10	354
HPV16		11	480		HPV16	E1	8	587
HPV16	E1	10	608		HPV16	E1	10	185
HPV16		11	530		HPV16	E1	11	185
HPV16	E1	10	593		HPV16	E1	9	289
HPV16	E1	9	512		HPV16	E1	10	289
HPV16	E1	10	512		HPV16	E1	8	253
HPV16	E1	8	561		HPV16	E1	9	253
HPV16	E1	9	190		HPV16	E1	10	253
HPV16	E1	10	553		HPV16	E1	8	525
HPV16	E1	11	553		HPV16	E1	8	585
HPV16	E1	11	302		HPV16	E1	10	585
HPV16	E1	10	600		HPV16	E1	11	498
HPV16	E1	10	577		HPV16		11	85
HPV16	E1	8	441		HPV16		11	197
HPV16	E1	8	556		HPV16		11	345
HPV16	E1	9	556		HPV16		11	443
HPV16	E1	8	419		HPV16	E1	10	24
HPV16	E1	10	419		HPV16	E1	9	584
HPV16		11	359		HPV16	E1	11	584
HPV16	E1	8	188		HPV16		8	274
HPV16		11	188		HPV16	E1	9	601

Table X HLA A24 Supermotif Peptides

				THE TEXT OUPCING	an replaces		
HPV16	E1	8	332		HPV16 E2	10	356
HPV16		9	332		HPV16 E2	11	288
HPV16		8	339		HPV16 E2	10	332
HPV16		10	509		HPV16 E2	11	82
HPV16		9	321		HPV16 E2	8	42
HPV16		10	531		HPV16 E2	10	42
			261		HPV16 E2	9	354
HPV16		11	578		HPV16 E2	10	354
HPV16		9	578		HPV16 E2	9	91
HPV16		11			HPV16 E2	8	177
HPV16		9	90		HPV16 E2	8	103
HPV16		10	320		HPV16 E2	11	16
HPV16		10	270				
HPV16		8	72		HPV16 E2 HPV16 E2	11 9	77 311
HPV16		11	72				
HPV16		11	331		HPV16 E2 HPV16 E2	11	311 157
HPV16		9	41			8	
HPV16		11	41		HPV16 E2	11	157
HPV16		11	228		HPV16 E2	8	296
HPV16		9	69		HPV16 E2	10	296
HPV16		11	69		HPV16 E2	8	127
HPV16		9	221		HPV16 E2	11	127
HPV16		8	63		HPV16 E2	8	284
HPV16		11	63		HPV16 E2	8	9
HPV16		8	314		HPV16 E2	11	9
HPV16		11	309		HPV16 E2	8	325
HPV16		8	124		HPV16 E2	11	325
HPV16		11	124		HPV16 E2	10	106
HPV16		8	25		HPV16 E2	11	60
HPV16		9	25		HPV16 E2	10	120
HPV16		11	246		HPV16 E2	9	170
HPV16		8	96		HPV16 E2	8	345
HPV16		8	31		HPV16 E2	8	76
HPV16		9	74		HPV16 E2	8	151
HPV16		10	74		HPV16 E2	9	151
HPV16		8	80		HPV16 E2	10	191
HPV16		9	185		HPV16 E2	8	349
HPV16		10	185		HPV16 E2	9	86
HPV16		8	118		HPV16 E2	8	304
HPV16		8	204		HPV16 E2	9	304
HPV16		11	100		HPV16 E2	11	278
HPV16		8	340		HPV16 E2	8	37
HPV16		11	346		HPV16 E2	9	7
HPV16		11	168		HPV16 E2	10	7
HPV16		9	163		HPV16 E2	10	302
HPV16		9	156		HPV16 E2	11	302
HPV16		9	230		HPV16 E2	8	23
HPV16		8	114		HPV16 E2	10	23
HPV16		8	29		HPV16 E2	11	23
HPV16		10	29		HPV16 E2	8	261
HPV16		10	53		HPV16 E2	11	261
HPV16		9	290		HPV16 E2	11	144
HPV16		8	35		HPV16 E2	8	355
HPV16		9	35		HPV16 E2	9	355
HPV16	E2	10	35		HPV16 E2	11	355
HPV16		9	18		HPV16 E2	10	61
HPV16	E2	8	130		HPV16 E2	10	78
HPV16	E2	9	130		HPV16 E2	9	297
HPV16		10	130		HPV16 E2	10	93
HPV16		8	193		HPV16 E2	11	93
HPV16	E2	8	356		HPV16 E2	8	334

Table X HLA A24 Supermotif Peptides

			Cupcinio	in replaces		
HPV16	E2	10	310	HPV16 E5	8	66
HPV16	E2	10	128	HPV16 E5	9	66
HPV16	E2	11	128	HPV16 E5	8	75
HPV16	E2	10	116	HPV16 E5	8	65
HPV16	E2	9	357	HPV16 E5	9	65
HPV16	E2	9	146	HPV16 E5	10	65
HPV16	E2	10	146	HPV16 E5	8	64
HPV16	E2	11	336	HPV16 E5	9	64
HPV16	E2	9	192	HPV16 E5	10	64
HPV16	E2	9	333 .	HPV16 E5	11	64
HPV16	E2	10	145	HPV16 E5	8	43
HPV16	E2	11	145	HPV16 E5	9	43
HPV16		8	147	HPV16 E5	8	44
HPV16		9	147	HPV16 E5	10	51
HPV16		11	147	HPV16 E5	11	51
HPV16		11	183	HPV16 E5	8	61
HPV16		10	101	HPV16 E5	9	61
HPV16		8	92 .	HPV16 E5	11	61
HPV16		11	92	HPV16 E5	10	71
HPV16		9	102	HPV16 E5	11	71
HPV16		8	312	HPV16 E5	8	73
HPV16		10	312	HPV16 E5	9	73
HPV16		8	131	HPV16 E5	10	73
HPV16		9	131	HPV16 E5	8	42
HPV16		11	115	HPV16 E5	9	42
HPV16		9	159	HPV16 E5	10	42
HPV16		10	159	HPV16 E5	9	11
HPV16		11	32	HPV16 E5	8	16
HPV16		11	154	HPV16 E5	8	22
HPV16		9	43	HPV16 E5	11	22
HPV16		10	158	HPV16 E5	8	27
	E2	11	158	HPV16 E5 HPV16 E5	8 11	32 32
HPV16		8	56	HPV16 E5 HPV16 E5	11	47
HPV16		9	56	HPV16 E5	10	33
HPV16		10	56	HPV16 E5	11	33
HPV16		11	56	HPV16 E5	10	48
HPV16		10	18	HPV16 E5	9	49
HPV16		11	18 59	HPV16 E5	11	1
HPV16 HPV16		10	59	HPV16 E5	9	3
HPV16		11	59	HPV16 E5	10	3
HPV16		9	14	HPV16 E5	11	70
HPV16		10	14	HPV16 E5	9	31
HPV16		8	26	HPV16 E5	8	41
HPV16		9	26	HPV16 E5	ء و	41
HPV16		9	24	HPV16 E5	10	41
HPV16		10	24	HPV16 E5	11	41
HPV16		11	24	HPV16 E5	8	8
HPV16		8	20	HPV16 E5	9	8
HPV16		9	20	HPV16 E5	10	8
HPV16		10	20	HPV16 E5	8	37
HPV16		8	5	HPV16 E5	9	37
HPV16		11	5	HPV16 E5	11	37
HPV16		9	60	HPV16 E5	8	35
HPV16		10	60	HPV16 E5	9	35
HPV16		9	72	HPV16 E5	10	35
HPV16	E5	10	72	HPV16 E5	11	35
HPV16	E5	11	72	HPV16 E5	8	10
HPV16	E5	8	15	HPV16 E5	10	10
HPV16		9	15	HPV16 E5	8	9

Table X HLA A24 Supermotif Peptides

				TILA A24 Supernio	in replices		
HPV16	E5	9	9		HPV16 E6	11	66
HPV16		11	9		HPV16 E6	10	21
HPV16		8	38		HPV16 E6	8	43
HPV16		10	38		HPV16 E6	10	43
HPV16		11	38		HPV16 E6	8	27
HPV16		8	21		HPV16 E6	9	27
HPV16		9	21		HPV16 E6	9	98
HPV16	E5	8	62		HPV16 E6	10	98
HPV16		10	62		HPV16 E6	11	98
HPV16		11	62		HPV16 E6	9	131
HPV16		8	67		HPV16 E6	8	151
HPV16		8	50		HPV16 E6	11	89
HPV16		11	50		HPV16 E6	11	29
HPV16		9	63		HPV16 E6	10	94
HPV16		10	63		HPV16 E6	8	28
HPV16	E5	11	63		HPV16 E6	11	93
HPV16	E5	9	39		HPV16 E6	8	38
HPV16		10	39 -		HPV16 E6	9	49
HPV16	E5	11	39		HPV16 E6	11	49
HPV16		9	53		HPV16 E6	8	60
HPV16	E6	9	68		HPV16 E7	9	68
HPV16		10	68		HPV16 E7	8	75
HPV16	E6	8	110		HPV16 E7	9	75
HPV16		10	58		HPV16 E7	10	75
HPV16	E6	8	73		HPV16 E7	9	81
HPV16		11	73		HPV16 E7	9	14
HPV16	E6	8	23		HPV16 E7	10	14
HPV16	E6	11	23		HPV16 E7	8	21
HPV16	E6	8	37		HPV16 E7	8	4
HPV16	E6	9	37		HPV16 E7	9	4
HPV16	E6	9	87		HPV16 E7	10	4
HPV16	E6	9	51		HPV16 E7	8	18
HPV16	E6	11	51		HPV16 E7	11	18
HPV16	E6	8	32		HPV16 E7	9	85
HPV16	E6	9	25		HPV16 E7	10	73
HPV16	E6	10	25		HPV16 E7	11	73
HPV16	E6	11	25		HPV16 E7	8	82
HPV16	E6	10	48		HPV16 E7	11	83
HPV16		9	82		HPV16 E7	11	12
HPV16		10	82		HPV16 E7	8	6
HPV16		8	76		HPV16 E7	10	6
HPV16		11	76		HPV16 E7	11	66
HPV16		8	92		HPV16 E7	8	77
HPV16		8	125		HPV16 E7	11	77
HPV16		11	125		HPV16 E7	9	71
HPV16		11	85		HPV16 E7	10	56
HPV16		11	34		HPV16 E7	10	78
HPV16		9	59		HPV16 E7	8	86 7
HPV16		9	75		HPV16 E7	9	
HPV16		8	79		HPV16 E7	10	19
HPV16		10	79		HPV16 E7	11	55 373
HPV16		9	18		HPV16 L1	9	292
HPV16		11	107		HPV16 L1 HPV16 L1	10 11	292
HPV16		9	44		HPV16 L1	9	70
HPV16		11	44		HPV16 L1	10	205
HPV16		10	90		HPV16 L1	11	172
HPV16		11	134		HPV16 L1	9	183
HPV16		10	102 116		HPV16 L1	10	251
HPV16		11 9	116		HPV16 L1	11	251
HPV16	oa.	9	0.0			11	

Table X HLA A24 Supermotif Peptides

				The true to apprint	. r optioo	•		
HPV16	L1	9	13		HPV16	L1	10	22
HPV16	L1	11	13		HPV16		8	102
HPV16	L1	9	249		HPV16		10	102
HPV16		11	484		HPV16		11	457
HPV16		8	397		HPV16		8	23
HPV16		11	397		HPV16		9.	23
HPV16		10	225		HPV16		8	332
HPV16		8	154		HPV16		9	401
HPV16		9	228		HPV16		9	424
HPV16		8	361		HPV16		9	86
HPV16		10	361		HPV16		11	86
HPV16		11	442		HPV16		11	11
HPV16		9	412		HPV16		8	407
HPV16		8	17		HPV16		10	407
HPV16		9	17		HPV16		11	407
HPV16		11	17		HPV16		9	6
HPV16		10	259		HPV16		10	59
HPV16		9	176		HPV16		11	59
HPV16		10	176		HPV16		9 10	108
HPV16		9	378		HPV16 HPV16			108
HPV16		8	132		HPV16		9	493 480
HPV16		8	474 245		HPV16		11	480
HPV16		10	395		HPV16		10	85
HPV16		8			HPV16		9	406
HPV16		10 9	395 388		HPV16		11	406
HPV16		11	388		HPV16		11	151
HPV16		8	52		HPV16		11	262
HPV16		9	52		HPV16		10	90
HPV16		10	52		HPV16		8	46
HPV16		8	24		HPV16		8	256
HPV16		9	273		HPV16		9	469
HPV16		10	273		HPV16		10	272
HPV16		8	400		HPV16		11	272
HPV16		10	400		HPV16	L1	8	184
HPV16		10	5		HPV16	L1	11	216
HPV16		10	472		HPV16	L1	8	68
HPV16		8	274		HPV16	L1	9	68
HPV16		9	274		HPV16	L1	11	68
HPV16	L1	9	116		HPV16	L1	8	409
HPV16		11	116		HPV16	L1	9	409
HPV16	L1	10	230		HPV16	L1	8	87
HPV16	L1	8	110		HPV16	L1	10	87
HPV16	L1	8	348		HPV16		10	29
HPV16	L1	9	348		HPV16		11	29
HPV16	L1	8	142		HPV16		11	414
HPV16	L1	11	142		HPV16		9	226
HPV16		8	499		HPV16		11	226
HPV16		10	499		HPV16		10	263
HPV16		10	431		HPV16		11	263
HPV16		9	93		HPV16		8	325
HPV16		11	93		HPV16		9	325
HPV16		9	438		HPV16		11	58
HPV16		8	166		HPV16		8	311
HPV16		10	166		HPV16		10	476
HPV16		10	130			L1	8	367
HPV16		9	140		HPV16		8	383
HPV16		10	140		HPV16		9	218
HPV16		8	22			L1	8	296
HPV16	L1	9	22		HPV16	пŢ	9	19

Table X HLA A24 Supermotif Peptides

				-	-			
HPV16	L1	11	19		HPV16	Ll	11	379
HPV16	L1	10	316		HPV16		8	219
HPV16		10	77		HPV16		11	358
HPV16		11	77		HPV16		10	415
HPV16		8	247		HPV16		11	415
HPV16	L1	11	247		HPV16		10	380
HPV16		9	213		HPV16		11	380
HPV16		8	489		HPV16		10	443
HPV16		11	138		HPV16		8	413
HPV16		8	466		HPV16		9	3
HPV16		11	43		HPV16		8	326
HPV16		8	267		HPV16 HPV16		10	359 20
HPV16		9	267		HPV16		10	20
HPV16		10	267		HPV16		11	20
HPV16		9 11	399 399		HPV16		9	30
HPV16 HPV16		8	487		HPV16		10	30
HPV16		9	487		HPV16		9	317
HPV16		10	487		HPV16		9	416
HPV16		8	331		HPV16		10	416
HPV16		9	331		HPV16		11	302
HPV16		11	181		HPV16		9	260
HPV16		8	280		HPV16		8	7
HPV16		8	2		HPV16		8	389
HPV16		10	2		HPV16		10	389
HPV16		9	95		HPV16		8	275
HPV16		8	445		HPV16		8	470
HPV16		10	100		HPV16	L1	8	53
HPV16		9	67		HPV16	L1	9	53
HPV16		10	67		HPV16	L1	9	60
HPV16	L1	10	115		HPV16	L1	10	60
HPV16	L1	8	253		HPV16	L2	9	241
HPV16	L1	9	253		HPV16		10	241
HPV16	L1	11	253		HPV16	L2	10	356
HPV16	L1	11	271		HPV16		11	356
HPV16		11	28		HPV16		9	293
HPV16	L1	9	174		HPV16		11	293
HPV16		11	174		HPV16		11	256
HPV16		10	419		HPV16		8	282
HPV16		9	324		HPV16		11	329
HPV16		10	324		HPV16		8	445
HPV16		10	308		HPV16		9	445 445
HPV16		11	308		HPV16		10 11	445
HPV16		11	49		HPV16		11	31
HPV16		9	422 422		HPV16		10	285
		11	365		HPV16		8	367
HPV16 HPV16		11	521		HPV16		10	84
HPV16		8	4		HPV16		9	140
HPV16		11	4		HPV16		8	261
HPV16		11	471		HPV16		8	195
HPV16		8	423		HPV16		10	340
HPV16		10	423		HPV16		8	176
HPV16		8	439		HPV16		8	242
HPV16		11	238		HPV16		9	242
HPV16		9	408		HPV16		11	242
HPV16		10	408		HPV16		8	181
HPV16		10	522		HPV16		8	446
HPV16		9	362		HPV16		9	446
HPV16		8	379		HPV16	L2	10	446
	-		-					

Table X HLA A24 Supermotif Peptides

				HLA A24 Supermotit	Peptide	5		
HPV16	L2	8	259		HPV16	L2	10	397
HPV16		9	259		HPV16		9	208
HPV16		10	259		HPV16		10	192
HPV16		10	364		HPV16		11	192
HPV16		11	364		HPV16		8	401
HPV16		8	26		HPV16		10	401
HPV16		8	65		HPV16	L2	11	417
HPV16		9	65		HPV16		11	215
HPV16	L2	11	65		HPV16	L2	9	429
HPV16		11	76		HPV16	L2	11	429
HPV16	L2	9	52		HPV16	L2	8	409
HPV16		9	392		HPV16	L2	10	409
HPV16	L2	11 .	392		HPV16	L2	9	161
HPV16	L2	9	180		HPV16	L2	10	172
HPV16	L2	9	325		HPV16	L2	8	358
HPV16	L2	8	439		HPV16	L2	9	358
HPV16	L2	9	439		HPV16	L2	11	358
HPV16	L2	10	439		HPV16	L2	8	221.
HPV16		10	32		HPV16		9	97
HPV16	L2	10	45		HPV16		10	381
HPV16	L2	11	45		HPV16		9	463
HPV16		8	420		HPV16		8	44
HPV16		9	420		HPV16		11	44
HPV16		11	420		HPV16		8	342
HPV16		8	243		HPV16		11	310
HPV16		10	243		HPV16		9	234
HPV16		11	135		HPV16		10	234
HPV16		8	250		HPV16 HPV16		9	47 47
HPV16		11	250		HPV16		11	436
HPV16 HPV16		9	286 430		HPV16		8	305
HPV16		10	430		HPV16		11	461
HPV16		11	430		HPV16		9	298
HPV16		10	105		HPV16		10	9
HPV16		11	105		HPV16		11	9
HPV16		10	248		HPV16	L2	8	313
HPV16	L2	8	39		HPV16	L2	11	302
HPV16	L2	10	39		HPV16	L2	8	319
HPV16	L2	11	35	1	HPV16	L2	9	319
HPV16	L2	8	323		HPV16		10	319
HPV16	L2	11	323		HPV16		10	274
HPV16		8	236		HPV16		11	274
HPV16		11	427		HPV16		9	360
HPV16		9	249		HPV16		10	360
HPV16		8	294		HPV16		9	125
HPV16		10	294		HPV16		11	125
HPV16		8	108		HPV16		11	104
HPV16		9	410		HPV16 HPV16		9	107
HPV16		11	410 365		HPV16		9	50
HPV16		9 10			HPV16		11	50
HPV16		10	365 266		HPV16		8	138
HPV16		9	454		HPV16		9	138
HPV16		11	454		HPV16		11	138
HPV16		8	276		HPV16		8	189
HPV16		9	276		HPV16		10	189
HPV16		11	276		HPV16		9	331
HPV16		10	407		HPV16		11	331
HPV16		9	419		HPV16		8	186
HPV16	L2	10	419	1	HPV16	L2	11	186

Table X HLA A24 Supermotif Peptides

				-				
HPV16	L2	8	387		HPV18		9	273
HPV16	L2	11	347		HPV18		10	273
HPV16	L2	10	384		HPV18		11	273
HPV16	L2	11	384		HPV18		9	311
HPV16	L2	8	162		HPV18		8	240
HPV16	L2	9	40		HPV18		9	240
HPV16	L2 .	8	332		HPV18		9	196
HPV16	L2	10	332		HPV18		10	196
HPV16	L2	9	438		HPV18		8	535
HPV16	L2	10	438		HPV18		9	535
HPV16	L2	11	438		HPV18		9	49
HPV16		8	399		HPV18		8	363
HPV16		10	399		HPV18		10	363
HPV16		10	187		HPV18		8	381
HPV16		9	85		HPV18		9	381
HPV16		10	311		HPV18		10	637
HPV16		11	265		HPV18		8	106
HPV16		8	156		HPV18		11	106
HPV16		9	398		HPV18		9	42
HPV16		11	398		HPV18		10	342
HPV16		8	141		HPV18 HPV18		9 11	342
HPV16		9	244		HPV18		10	220
HPV16		11	351 136		HPV18		8	540
HPV16 HPV16		10 11	136		HPV18		8	445
HPV16		8	350		HPV18		9	445
HPV16		11	153		HPV18		11	445
HPV16		8	209		HPV18		9	459
HPV16		10	154		HPV18		8	594-
HPV16		10	251		HPV18	E1	10	610
HPV16		10	348		HPV18	E1	11	610
HPV16	L2	8	53		HPV18		8	62
HPV16	L2	9	155		HPV18		9	62
HPV16	L2	8	464		HPV18		9	108
HPV16	L2	9	267		HPV18		11	108
HPV16		11	267		HPV18		8	375
HPV16		8	393		HPV18		10	375
HPV16		10	393		HPV18		11	366
HPV16		8	447		HPV18		11 10	309 104
HPV16		9	447		HPV18		9	74
HPV16		10	453		HPV18		9	338
HPV16		10	437		HPV18		10	338
HPV16 HPV16		11 9	349		HPV18		8	345
HPV16		11	452		HPV18		9	345
HPV16		8	326		HPV18		9	619
HPV18		11	398		HPV18		11	619
HPV18		9	246		HPV18		8	50
HPV18		11	22		HPV18	E1	10	497
HPV18		9	546		HPV18	E1	10	265
HPV18		9	325		HPV18	E1	9	500
HPV18		10	213		HPV18		8	460
HPV18		11	526		HPV18		11	460
HPV18	E1	8	40		HPV18		8	463
HPV18	E1	11	40		HPV18		9	463
HPV18	E1	8	531		HPV18		11	470
HPV18		9	531		HPV18		10	399
HPV18		9	216		HPV18		11	399
HPV18		11	216		HPV18		9	452
HPV18	El	10	618		HPV18	EI	10	452

Table X HLA A24 Supermotif Peptides

				IILA A24 3	upermon re	pudes			
HPV18	E1	8	226		HP	V18	E1	11	304
HPV18		8	130			V18		11	204
HPV18		8	508			V18		11	285
HPV18		9	508			V18		9	376
HPV18		10	508			V18		11	376
HPV18		9	257		HP	V18	E1	8	520
HPV18		10	257			V18		9	520
HPV18		11	257			V18		8	350
HPV18		9	356		HP	V18	E1	10	295
HPV18		9	332		HP	V18	E1	11	295
HPV18		11	223		HP	V18	E1	8	267
HPV18		8	494		HP	V18	E1	8	326
HPV18		9	55		HP	V18	E1	11	326
HPV18		8	11		HP	V18	E1	9	214
HPV18		9	11		HP	V18	E1	11	214
HPV18		8	473		HP	V18	E1	10	527
HPV18		10	473		HP	V18	E1	8	312
HPV18		8	182		HP	V18	E1	11	47
HPV18		8	279		HP	V18	E1	10	367
HPV18	E1	10	279		HP	V18	E1	10	361
HPV18		10	249		HP	V18	E1	11	188
HPV18		8	16		HP	V18	E1	10	574
HPV18		8	499		HP	V18	E1	8	428
HPV18	E1	10	499		HP	V18	E1	9	487
HPV18	E1	9	270		HP	V18	E1	11	487
HPV18	E1	8	306		HP	V18	E1	11	158
HPV18	E1	9	306			V18		8	191
HPV18	E1	10	418			V18		11	191
HPV18	E1	8	247			V18^		10	615
HPV18	E1	11	352			V18		10	600
HPV18	E1	9	266				E1	8	264
HPV18	E1	10	461				E1	11	264
HPV18	E1	11	461			V18		8	568
HPV18	E1	10	590			V18		11	551
HPV18		10	23			V18		8	448
HPV18	E1	10	449			V18		11	448
HPV18		8	439			V18		10	560
HPV18		11	439			V18		11	560
HPV18		11	647			V18		9	519
HPV18		9	318			V18		10	519 252
HPV18		10	318			V18		10	252
HPV18		11	318			V18		11	607
HPV18		8	259			V18		10	584
HPV18		9	259				E1	8	451
HPV18		10	259 237				E1	10	451
HPV18		11					E1	11	451
HPV18		9	206				E1	8	457
HPV18 HPV18		10	277				E1	11	457
HPV18		9 10	277				E1	8	563
HPV18		9	407			V18		9	563
		9	129			V18		8	426
HPV18		8	281			V18		10	426
HPV18		9	561			V18		10	80
HPV18		10	561			V18		8	369
HPV18		11	561			V18		10	431
HPV18		8	261			V18		11	589
	E1	11	261				E1	8	102
HPV18		9	304				E1	10	128
HPV18		10	304			V18		9	349
111 410									

Table X HLA A24 Supermotif Peptides

						· · · · · · · · · · · · · · · · · · ·			
HPV18	E1	8	294		3	HPV18	El	8	536
HPV18	El	11	294		3	HPV18	E1	9	328
HPV18	E1	11	330		1	HPV18	El	10	538
HPV18	E1	8	622		1	HPV18	E1	9	492
HPV18	E1	9	582		1	HPV18	E1	10	492
HPV18	E1	9	287		1	HPV18	E1	9	585
HPV18	E1	8	454		1	HPV18	E1	11	585
HPV18		10	454		1	HPV18	E1	8	408
HPV18		11	454		3	HPV18	E1	11	542
HPV18		11	496		1	HPV18	E1	10	327
HPV18		9	225		1	HPV18	E2	8	76
HPV18		9	507		1	HPV18	E2	11	76
HPV18		10	507		1	HPV18	E2	11	45
HPV18		11	507		1	HPV18	E2	8	351
HPV18		9	553		1	HPV18	E2	9	351
HPV18		10	553		1	HPV18	E2	10	82
HPV18		11	93		1	HPV18	E2	9	154
HPV18	E1	11	302		1	HPV18	E2	10	154
HPV18		10	511		1	HPV18	E2	11	154
HPV18		8	322		I	HPV18	E2	8	214
HPV18		11	179		1	HPV18	E2	9	246
HPV18		8	491		1	HPV18	E2	8	137
HPV18		10	491		1	HPV18	E2	11	132
HPV18		11	491		1	HPV18	E2	10	14
HPV18		8	56		1	HPV18	E2	8	156
HPV18		9	462		1	HPV18	E2	9	156
HPV18		10	462		I	HPV18	E2	8	29
HPV18		9	253		F	HPV18	E2	9	29
HPV18		10	253	-	F	HPV18	E2	11	29
HPV18	E1	8	197		F	IPV18	E2	8	315
HPV18		9	197		I	HPV18	E2	8	210
HPV18	E1	8	260		F	HPV18	E2	11	95
HPV18		9	260		Ŧ	HPV18	E2	9	175
HPV18	E1	10	303		I	HPV18	E2	9	78
HPV18	E1	11	303		I	HPV18	E2	11	104
HPV18	E1	10	238			HPV18		8	340
HPV18	E1	11	238		F	IPV18	E2	9	190
HPV18	E1	10 .	533				E2	9	47
HPV18	E1	11	533			HPV18		9	161
HPV18	E1	8	532			IPV18		9	168
HPV18	E1	11	532			HPV18		9	291
HPV18	E1	9	296			HPV18		9	22
HPV18	E1	10	296			HPV18		9	312
HPV18	E1	9	591			HPV18		11	312
HPV18		11	591			IPV18		10	46
HPV18		11	537			HPV18		11	289
HPV18		9	221			IPV18		8	358
HPV18		8	592			IPV18		8	345
HPV18		10	592			IPV18		10	280
HPV18		8	217			IPV18		11	280
HPV18		10	217			IPV18		11	152
HPV18		11	505			IPV18		9	329
HPV18		9	81			IPV18		8	182
HPV18		9	280			IPV18		8	39
HPV18		8	339			IPV18		9	39
HPV18		9	339			IPV18		10	39
HPV18		11	244			IPV18		10	105
HPV18		9	608			IPV18		8	162
HPV18		8	346			IPV18		11	162
HPV18	E1	9	432		1	IPV18	EZ	10	133

Table X HLA A24 Supermotif Peptides

			HLA A24 Supermon	r Pepudes		
HPV18	E2	11	133	HPV18 E2	11	36
HPV18		11	67	HPV18 E2	9	142
HPV18	E2	8	107	HPV18 E2	10	163
HPV18	E2	9	185	HPV18 E2	11	163
HPV18		8	285	HPV18 E5	9	49
HPV18	E2	11	348	HPV18 E5	10	49
HPV18	E2	9	196	HPV18 E5	11	49
HPV18	E2	10	272	HPV18 E5	9	47
HPV18	E2	9	357	HPV18 E5	11	47
HPV18	E2	8	33	HPV18 E5	8	32
HPV18	E2	10	33	HPV18 E5	9	32
HPV18		9	38	HPV18 E5	8	30
HPV18		10	38	HPV18 E5	10	30
HPV18		11	38	HPV18 E5	11	30
HPV18		9	220	HPV18 E5	8	56
HPV18		11	56	HPV18 E5 HPV18 E5	9 11	56 56
HPV18		11	2	HPV18 E5	9	27
HPV18		8	61	HPV18 E5	11	27
HPV18		9	35 305	HPV18 E5	10	13
HPV18		9	11	HPV18 E5	11	13
HPV18 HPV18		10	11	HPV18 E5	9	6
HPV18		10	343	HPV18 ES	10	6
HPV18		8	41	HPV18 E5	8	57
HPV18		9	90	HPV18 E5	10	57
HPV18		11	303	HPV18 E5	8	50
HPV18		9	298	HPV18 E5	9	50
HPV18		11	230	HPV18 E5	10	50
HPV18		8-	233	HPV18 E5	8	65
HPV18		8	355	HPV18 E5	10	19
HPV18		9	355	HPV18 E5	10	5
HPV18		11	355	HPV18 E5	11	5
HPV18	E2	11	140	HPV18 E5	8	43
HPV18	E2	10	57	HPV18 E5	11	43
HPV18	E2	9	97	HPV18 E5	11	40
HPV18	E2	10	97	HPV18 E5	8	7
HPV18	E2	10	349	HPV18 E5	9	7
HPV18		11	349	HPV18 E5	11	4
HPV18		11	211	HPV18 E5	8	63
HPV18		10	231	HPV18 E5	10 8	63 62
HPV18		11	188	HPV18 E5	9	62
HPV18		11	336	HPV18 E5 HPV18 E5	11	62
HPV18		9	134 134	HPV18 E5	9	58
HPV18		10 11	134	HPV18 E5	10	22
HPV18 HPV18		8	197	HPV18 E5	9	35
HPV18		11	123	HPV18 E5	9	61
HPV18		10	141	HPV18 E5	10	61
HPV18		11	322	HPV18 E5	8	1
HPV18		10	96	HPV18 E5	10	1
HPV18		11	96	HPV18 E5	9	14
HPV18		10	124	HPV18 E5	10	14
HPV18		11	173	HPV18 E5	8	21
HPV18		8	143	HPV18 E5	11	21
HPV18		8	135	HPV18 E5	10	60
HPV18		9	135	HPV18 E5	11	60
HPV18		10	135	HPV18 E5	8	3
HPV18		9	164	HPV18 E5	9	25
HPV18	E2	10	164	HPV18 E5	11	25
HPV18	E2	11	164	HPV18 E5	8	51

PCT/US00/33549

Table X HLA A24 Supermotif Peptides

				•			
HPV18	E5	9	51	HPV18 E	6	9	93
HPV18	E5	11	51	HPV18 E	6	10	93
HPV18	E5	8	54	HPV18 E	6	11	93
HPV18		9	54	HPV18 E		9	98
HPV18		10	54	HPV18 E		8	95
HPV18		11	54	HPV18 E		9	95
HPV18		8	36	HPV18 E		9	22
HPV18		9	42	HPV18 E		8	111
HPV18		10	34	HPV18 E		11	111
HPV18		10	41	HPV18 E		8	7
HPV18		8	52	HPV18 E		11	7
HPV18		10	52	HPV18 E		11	11
HPV18		11	52	HPV18 E		10	3
HPV18		8	33	HPV18 E		9	126
HPV18		11	33	HPV18 E		8	74
HPV18		8	15	HPV18 E		11	59
HPV18		9	15	HPV18 E		11	24
HPV18		9	53 .	HPV18 E		10	84
HPV18		10	53	HPV18 E		10	89
HPV18	E5	11	53	HPV18 E		11	89
HPV18		9	48	HPV18 E		9 11	37 37
HPV18		8	68	HPV18 E		9	
HPV18		11	68	HPV18 E			44
HPV18 HPV18		8	105 18	HPV18 E			38
HPV18		11	18	HPV18 E			38
HPV18		8	32	HPV18 E			33
HPV18		10	32	HPV18 E			85
HPV18	E6	9	70	HPV18 E			85
HPV18		8	27	HPV18 E			6
HPV18		10	16	HPV18 E			6
HPV18		10	51	HPV18 E			63
	E6	11	88	HPV18 E			24
	E6	8	46	HPV18 E			82
HPV18		11	46	HPV18 E			82
HPV18		11	29	HPV18 E	7 :	10	82
HPV18		9	20	HPV18 E	7 :	10	40
	E6	11	20	HPV18 E	7 :	11	90
HPV18	E6	10	77	HPV18 E	7 8	В	14
HPV18	E6	8	40	HPV18 ET	7 :	11	11
HPV18	E6	10	40	HPV18 ET	7 :	11	73
HPV18		10	43	HPV18 E7	7 1	В	89
HPV18	E6	11	43	HPV18 E7	7 :	10	74
HPV18	E6	8	53	HPV18 E7			92
HPV18	E6	8	71	HPV18 E7		10	22
HPV18		11	71	HPV18 E7			88
	E6	10	97	HPV18 E7			7
HPV18	E6	8	120	HPV18 E7			93
	E6	11	120	HPV18 E7			12
	E6	11	80	HPV18 E7			75
HPV18		10	30	HPV18 L1			63
	E6	9	13	HPV18 L1			128
	E6	11	117	HPV18 L1			218
	E6	8	92	HPV18 L1			310
	E6	10	92	HPV18 L1			2
	E6	11	92	HPV18 L1			2
	E6	10	36	HPV18 L1			2
	E6	9	52	HPV18 L1			441
HPV18		11	102	HPV18 L1			350
HPV18	₽6	9	41	HPV18 L1	. 9	•	284

271

Table X HLA A24 Supermotif Peptides

				HLA A24 Supermout Fe	pudes			
HPV18	L1	8	122	HP	V18	L1	9	428
HPV18		10	122	HP	V18	L1	10	428
HPV18	Ll	11	520	HP	V18	L1	11	428
HPV18	L1	8	512	HP	V18	L1	8	11
HPV18	L1	10	512			L1	11	11
HPV18	L1	8	433	HP	V18	L1	8	58
HPV18	L1	11	433	HP	V18	L1	9	437
HPV18	L1	10	260			Ll	10	94
HPV18	L1	9	263			Ll	11	94
HPV18	L1	8	276			L1	8	40
HPV18	L1	10	276			Ll	9	40
HPV18	L1	8	396			L1	10	40
HPV18		10	396		V18		11	40
HPV18		8	330			L1	8	39
HPV18		9	330			L1	9	39
HPV18	L1	11	478			L1	10	39
HPV18		9	448			L1	11	39
HPV18		10	203		V18		11	46
HPV18		9	211		V18		10	47
HPV18		10	211		V18		8	219
HPV18		10	294			L1	9	9
HPV18		8	87			L1	10	9
HPV18		9	87			L1	8	32
HPV18		10	87		V18		9	32
HPV18		8	167		V18		10	32
HPV18		10	280			L1		443
HPV18		8	431		V18			443
HPV18		10	431					443
HPV18		9	308			L1		360
HPV18		10	308			L1	9	360 151
HPV18		8	436		V18 V18		9 11	151
HPV18		10	436			L1	9	143
HPV18		8	49			L1	10	143
HPV18		10	49				8	529
HPV18		11	321		V18		9	516
HPV18 HPV18		10	508				11	516
HPV18		9	95			L1		507
HPV18		10	95		V18		9	505
HPV18		8	145		V18		10	125
HPV18		8	535		V18			291
HPV18		8	177		V18			48
HPV18		11	177		V18		11	48
HPV18		11	342	HP	V18	L1	8	367
HPV18		9	233		V18		8	8
HPV18		11	326		V18		10	8
HPV18		8	383	HP	V18	L1	11	8
HPV18		9	383	HP	V18	L1	8	14
HPV18		10	165	HP	V18	Ll	9	14
HPV18	L1	9	175	HP	V18	L1	8	103
HPV18		10	175	HP	V18	L1	9	103
HPV18	L1	10	467	HP	V18	L1		445
HPV18	L1	10	265		V18		9	445
HPV18	L1	9	320	HP	V18	L1	8	104
HPV18		8	38	HP	V18	L1	10	159
HPV18	L1	9	38		V18		8	33
HPV18	L1	10	38		V18		9	33
HPV18		11	38		V18		10	64
HPV18		9	13		V18		11	64
HPV18	L1	10	13	. HP	V18	L1	11	17

Table X HLA A24 Supermotif Peptides

				III. I III. I Dapaimoni	· r cpmao			
HPV18	L1	9	21		HPV18	L1	10	6
HPV18	L1	8	336		HPV18	L1	10	135
HPV18	L1	8	3		HPV18	L1	8	81
HPV18	L1	10	3		HPV18	L1	8	288
HPV18		10	307		HPV18	L1	9	288
HPV18		11	307		HPV18	L1	11	288
HPV18		9	261		HPV18		11	93
HPV18		11	261		HPV18		8	459
HPV18		10	36		HPV18	L1	9	31
HPV18		11	36		HPV18		10	31
HPV18		11	84		HPV18	L1	11	31
HPV18		9	253		HPV18		9	359
HPV18		8	510		HPV18	L1	10	359
HPV18		10	510		HPV18		10	150
HPV18		9	54		HPV18		9	518
HPV18		11	54		HPV18		8	475
HPV18		8	52		HPV18	L1	9	335
HPV18		9	52		HPV18		11	306
HPV18		11	52		HPV18		10	455
HPV18		11	207		HPV18		10	408
HPV18		8	57		HPV18		11	78
HPV18		9	57		HPV18		9	209
HPV18		8	282		HPV18		11	209
HPV18		11	282		HPV18		10	451
HPV18		9	248		HPV18		11	451
HPV18		11	173		HPV18		9	442
HPV18		8	28		HPV18		11	442
HPV18		9	28		HPV18		11	273
HPV18		8	26		HPV18		9	444
HPV18		10	26		HPV18	L1	10	444
HPV18		11	26		HPV18 HPV18	L1	10	327
HPV18		10	240		HPV18	L1	11	327
HPV18		8	20		HPV18		9	397
HPV18		10	20		HPV18		8	473
HPV18		9	472		HPV18		10	473
HPV18		11	472		HPV18	L1	8	254 -
HPV18		11	412		HPV18		8	331
HPV18		8	502		HPV18	L1	11	393
HPV18		8	302		HPV18		10	479
HPV18		10	302		HPV18	L1	8	55
HPV18		9	435		HPV18	L1	10	55
HPV18		11	435		HPV18		11	55
HPV18		11	216		HPV18	L1	8	7
HPV18	L1	8	315		HPV18	L1	9	7
HPV18	L1	8	366		HPV18	L1	11	7
HPV18		9	366		HPV18	L1	8	449
HPV18	L1	8	137		HPV18		8	89
HPV18	L1	10	137		HPV18	L1	8	361
HPV18	L1	11	297		HPV18		10	394
HPV18	L1	10	416		HPV18	Ll	9	160
HPV18	L1	8	523		HPV18		8	34
HPV18	L1	9	523		HPV18		10	351
HPV18		9	130		HPV18		9	452
HPV18	L1	9	424		HPV18		10	452
HPV18		11	424		HPV18		9	295
HPV18	L1	8	481		HPV18	L1	11	35
HPV18	L1	9	102		HPV18		9	4
HPV18		10	102		HPV18		11	4
HPV18		11	158		HPV18		8	88
HPV18		9	6 .		HPV18	Ll	9	88

Table X
HLA A24 Supermotif Peptides

				ILA A24 Supermon	repude	,		
HPV18	L2	9	255		HPV18	L2	8	52
HPV18		11	255		HPV18		9	279
HPV18		9	370		HPV18		9	44
HPV18		8	161		HPV18		10	44
HPV18		11	286		HPV18		11	4.4
HPV18		8	341		HPV18		10	405
HPV18		10	341		HPV18		10	143
HPV18		11	341		HPV18		8	249
HPV18		11	303		HPV18		8	43
HPV18		8	275		HPV18		10	43
HPV18		10	278		HPV18		11	43
HPV18		11	322		HPV18		11	34
HPV18			404		HPV18		9	347
		11	142		HPV18		9	248
HPV18 HPV18		11 10	346		HPV18		8	242
			83		HPV18		10	242
HPV18		11	270		HPV18		10	287
HPV18					HPV18		10	391
HPV18		10	270		HPV18		10	338
HPV18		11	270					338
HPV18		10	396		HPV18		11	
HPV18		11	30		HPV18		8	437
HPV18		9	240		HPV18		9	305
HPV18		10	240		HPV18		10	386
HPV18		8	331		HPV18		8	325
HPV18		8	371		HPV18		9	325
HPV18		11	443		HPV18		11	325
HPV18		8	241		HPV18		11	390
HPV18		9	241		HPV18		11	337
HPV18		11	241		HPV18		9	419
HPV18		9	122		HPV18		10	419
HPV18		11	157		HPV18		9	98
HPV18		8	306		HPV18		10	98
HPV18		8	319		HPV18		9	120
HPV18		9	51		HPV18		11	120
HPV18		8	429		HPV18		9	376
HPV18	L2	9	429		HPV18		8 -	86
HPV18		10	429		HPV18		8	185
HPV18		8	64		HPV18		11	185
HPV18		9	64		HPV18		11	216
HPV18		11	64		HPV18		8	258
HPV18		8	188		HPV18		9	95
HPV18		10	188		HPV18		10	360
HPV18		9	432		HPV18		8	398
HPV18		10	432		HPV18		10	398
HPV18	L2	11	432		HPV18		8	452
HPV18		10	183		HPV18		8	312
HPV18	L2	10	310		HPV18		9	312
HPV18	L2	11	310		HPV18		10	172
HPV18	L2	8	37		HPV18		9	233
HPV18	L2	9	37		HPV18		10	233
HPV18		11	37		HPV18		8	46
HPV18	L2	8	134		HPV18		9	46
HPV18	L2	10	134		HPV18		10	426
HPV18	L2	8	292		HPV18		11	426
HPV18	L2	10	191		HPV18		11	295
HPV18	L2	9	318		HPV18		8	298
HPV18	L2	8	434		HPV18		9	298
HPV18	L2	9	434		HPV18		8	220
HPV18	L2	10	434		HPV18		8	316
HPV18	L2	11	434		HPV18	L2	11	316

Table X HLA A24 Supermotif Peptides

				TILA A24 Supermon Fej	pudes		
HPV18	T-2	10	450	HP	V31 E1	10	620
HPV18		11	368		V31 E1	8	508
HPV18		9	49		V31 E1	9	508
HPV18		11	49		V31 E1	9	49
HPV18		10	247		V31 E1	8	96
HPV18		11	147		V31 E1	10	96
HPV18		10	153		V31 E1	8	421
HPV18		8	365		V31 E1	11	421
HPV18		10	365		V31 E1	8	336
HPV18		8	409		V31 E1	10	336
HPV18		8	235		V31 E1	9	42
HPV18		9	149		V31 E1	10	42
HPV18		9	383		V31 E1	9	332
HPV18		8	121		V31 E1	8	528
HPV18		10	121		V31 E1	10	528
HPV18		8	377		V31 E1	8	348
HPV18		9	39		V31 E1	10	348
HPV18		9	406		V31 E1	9	311
HPV18		11	406		V31 E1	10	311
HPV18		10	304		V31 E1	8	418
HPV18		8	38		V31 E1	9	418
HPV18		10	38		V31 E1	11	418
HPV18		9	154		V31 E1	8	102
HPV18		8	136		V31 E1	11	102
HPV18		9	366		V31 E1	9	432
HPV18		9	135		V31 E1	11	432
HPV18		8	388		V31 E1	9	354
HPV18		10	217		V31 E1	10	583
HPV18		11	217		V31 E1	11	583
HPV18		10	113		V31 E1	8	193
HPV18		9	387		V31 E1	10	193
HPV18		10	31		V31 E1	11	193
HPV18		11	112		V31 E1	8	168
HPV18		9	399		V31 E1	11	168
HPV18		9	427		V31 E1	9	318
HPV18		10	427		V31 E1	9	592
HPV18		11	427		V31 E1	11	592
HPV18		8	436		V31 E1	8	50
HPV18		9	436		V31 E1	11	443
HPV18		8	400		V31 E1	10	372
HPV18		8	435		V31 E1	9	473
HPV18		9	435		V31 E1	9	425
HPV18		10	435		V31 E1	8	436
HPV31		11	371	HP	V31 E1	9	436
HPV31		9	519	HP7	V31 E1	8	199
HPV31		10	533	HP7	V31 E1	9	566
HPV31		11	533	HPV	V31 E1	8	433
HPV31	E1	9	298	HP/	V31 E1	10	433
HPV31	E1	9	186	HPV	V31 E1	11	433
HPV31		10	186	HP7	V31 E1	11	488
HPV31	E1	8	504	HP7	V31 E1	9	230
HPV31		9	504	HP7	V31 E1	11	230
HPV31		11	22	HP/	V31 E1	11	499
HPV31		9	81		V31 E1	8	467
HPV31		8	279		V31 E1	9	305
HPV31		9	279	HPV	V31 E1	10	252
HPV31		9	239	HPV	V31 E1	8	11
HPV31		9	284	HPV	V31 E1	9	11
HPV31		9	213	HPV	/31 E1	10	225
HPV31		10	100	HPV	V31 E1	11	225

Table X HLA A24 Supermotif Peptides

				***************************************	o apermoni	replice	•		
HPV31	E1	8	446			HPV31	E1	9	253
HPV31	E1	9	446			HPV31	E1	11	143
HPV31	E1	10	446			HPV31	E1	11	340
HPV31	E1	8	196			HPV31	E1	10	547
HPV31	E1	11	196			HPV31	E1	9	104
HPV31		10	222			HPV31		11	104
HPV31		9	16			HPV31		9	135
HPV31		9	243			HPV31		9	460
HPV31		11	243			HPV31		11	460
HPV31		11	510			HPV31		10	588
HPV31		10	391			HPV31		10	573
HPV31		11	478			HPV31			492
HPV31		10	268			HPV31		10	492
HPV31		11	268			HPV31		8	541
HPV31		8	381			HPV31		11	93
HPV31		10	422			HPV31		9	170
HPV31		10	190			HPV31		11	524
HPV31		11	190			HPV31		10	580
HPV31		8	424			HPV31		8	237
HPV31		10	424			HPV31		11	237
HPV31		9	380			HPV31		10	557
HPV31		10	276			HPV31		8	430
HPV31		11	276			HPV31		11	430
HPV31		9	272			HPV31		8	536
HPV31		9	291			HPV31		9	536
HPV31		10	291			HPV31		8	399
HPV31		11	291			HPV31		8	323
HPV31		8	119			HPV31		9	145
HPV31		10	119			HPV31		9	342
HPV31		9	232			HPV31		9	260
HPV31		10	232			HPV31		11	260
HPV31		9	179			HPV31		8	267
HPV31		10	179			HPV31		11	267
HPV31		8	412			HPV31		10	124
HPV31		11	412			HPV31		11	562
HPV31		10	88			HPV31		11	303
HPV31		9	125			HPV31		8	595
HPV31		9	470			HPV31		9	555
HPV31		10	470			HPV31		10	427
HPV31		9	277			HPV31		11	427
HPV31		10	277			HPV31		10	591
HPV31		11	277			HPV31		8	472
HPV31		8	273			HPV31		10	472
HPV31		8	234			HPV31		8	435
HPV31		11	234			HPV31		9	435
HPV31		9	534			HPV31		10	435
HPV31		10	534			HPV31		9	198
HPV31		11	534			HPV31		9	329
HPV31		11	258			HPV31		9	526
HPV31		10	563			HPV31		10	526
HPV31		8	240			HPV31	E1	8	246
HPV31		9	194			HPV31		10	246
HPV31		10	194			HPV31		11	246
HPV31		8	299			HPV31		10	469
HPV31		11	299			HPV31		11	469
HPV31		10	500			HPV31		8	294
HPV31		8	187			HPV31		9	294
HPV31		9	187			HPV31		11	211
HPV31		8	285			HPV31	E1	11	616
HPV31		11	47			HPV31		8	250

Table X HLA A24 Supermotif Peptides

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HPV31	E1	9	250		HPV31	E2	8	124
HPV31	E1	9	480		HPV31	E2	11	124
HPV31		10	480		HPV31	E2	9	91
HPV31	E1	11	480		HPV31	E2	11	91
HPV31	E1	8	464		HPV31	E2	8	31
HPV31		10	464		HPV31	E2 ·	11	204
HPV31	E1	11	464		HPV31	E2	9	74
HPV31	E1	10	334		HPV31	E2	11	100
HPV31	E1	10	617		HPV31	E2	8	80
HPV31	E1	8	567		HPV31	E2	9	185
HPV31	E1	9	269		HPV31	E2	11	185
HPV31		10	269		HPV31		9	86
HPV31	E1	8	233		HPV31		8	171
HPV31	E1	9	233		HPV31		11	168
HPV31	E1	8	333		HPV31		10	156
HPV31	E1	11	333		HPV31		8	114
HPV31	E1	8	505		HPV31		8	29
HPV31		11	505		HPV31		10	29
HPV31	E1	9	226		HPV31		8	35
HPV31		10	226		HPV31		9	35
HPV31		8	565		HPV31		10	35
HPV31		10	565		HPV31		8	164
HPV31		10	23		HPV31		8	297
HPV31		11	84		HPV31		9	297
HPV31		11	177		HPV31		9	18
HPV31		11	325		HPV31 HPV31		9	130 130
HPV31		9	349		HPV31		10	130
HPV31		11	349 254		HPV31		10	295
HPV31		9	223		HPV31		11	295
HPV31		9	564		HPV31		9	304
HPV31		11	564		HPV31		8	193
HPV31		9	581		HPV31		9	157
HPV31		8	312		HPV31		11	157
HPV31		9	312		HPV31		11	183
HPV31		8	17		HPV31	E2	8	177
HPV31		8	319		HPV31	E2	8	42
HPV31		10	489		HPV31	E2	10	42
HPV31		9	301		HPV31	E2	8	103
HPV31	E1	9	465		HPV31	E2	9	318
HPV31	E1	10	465		HPV31	E2	11	318
HPV31	E1	10	511		HPV31		10	101
HPV31	E1	9	558		HPV31		10	78
HPV31		11	558		HPV31		11	77
HPV31		9	89		HPV31		9	43
HPV31		10	300		HPV31		9	170
HPV31		8	72		HPV31		8	303
HPV31		11	72		HPV31		10	303
HPV31		8	338		HPV31		9	84 84
HPV31		9	69		HPV31 HPV31		11	254
HPV31		10	69		HPV31		9	254
HPV31		11	69		HPV31		8	127
HPV31		10 8	61 291		HPV31		10	127
HPV31		10	286		HPV31			127
HPV31		11	286		HPV31		9	361
HPV31		9	307		HPV31		10	361
HPV31		10	330		HPV31			9
HPV31		10	40		HPV31		11	9
HPV31		8	352		HPV31		9	133
		-						

Table X HLA A24 Supermotif Peptides

				TILIT TILY Supermoun	I optioo.			
HPV31	E2	11	294		HPV31	E5	9	40
HPV31	E2	10	106		HPV31	E5	10	40
HPV31		10	120	`	HPV31	E5	11	40
HPV31	E2	10	317		HPV31	E5	8	53
HPV31	E2	8	96		HPV31	E5	10	53
HPV31	E2	10	191		HPV31	E5	11	53
HPV31	E2	8	151		HPV31	E5	8	59
HPV31	E2	9	151		HPV31	E5	9	59
HPV31	E2	8	321		HPV31	E5	10	59
HPV31	E2	8	25		HPV31	E5	11	59
HPV31	E2	9	25		HPV31	E5	10	18
HPV31	E2	8	37		HPV31	E5	9	14
HPV31	E2	9	7		HPV31	E5	10	14
HPV31	E2	10	7		HPV31	E5	11	14
HPV31	E2	8	311		HPV31	E5	8	61
HPV31	E2	9	311		HPV31	E5	9	61
HPV31	E2	10	309		HPV31	E5	11	61
HPV31	E2	11	309		HPV31		9	26
HPV31	E2	9	206		HPV31	E5	8	20
HPV31		10	53		HPV31	E5	10	20
HPV31		8 .	346		HPV31	E5	9	3
HPV31	E2	9	346		HPV31	E5	11	3
HPV31		10	266		HPV31	E5	9	66
HPV31		8	198		HPV31	E5	8	15
HPV31		10	198		HPV31	E5	9	15
HPV31		8	63		HPV31	E5	10	15
HPV31		11	63		HPV31	E5	9	24
HPV31		9	364		HPV31	E5	11	24 72
HPV31		9	128		HPV31 HPV31	E5 E5	9 10	72
HPV31		10	128 128		HPV31	E5	10	48
HPV31 HPV31		11 9	93		HPV31	E5	9	11
HPV31		10	93		HPV31	E5	8	67
HPV31		11	93		HPV31	E5	8	62
HPV31		10	221		HPV31	E5	10	62
HPV31		11	220		HPV31	E5	11	62
HPV31		8	362		HPV31	E5	10	23
HPV31		9	362		HPV31	E5	10	71
HPV31		11	362		HPV31	E5	11	71
HPV31		11	343		HPV31	E5	8	45
HPV31		9	199		HPV31	E5	8	16
HPV31	E2	9	192		HPV31	E5	9	16
HPV31	E2	9	41		HPV31	E5	8	22
HPV31	E2	11	41		HPV31	E5	11	22
HPV31	E2	11	147		HPV31	E5	9	44
HPV31	E2	8	92		HPV31	E5	8	43
HPV31	E2	10	92			E5	10	43
HPV31	E2	11	92		HPV31	E5	8	42
HPV31		10	344		HPV31	E5	9	42
HPV31		11	344		HPV31	E5	11	42
HPV31		9	102		HPV31	E5	8	27
HPV31		8	131		HPV31	E5	8	32
	E2	9	131		HPV31	E5	11	32
HPV31	E2	11	131		HPV31	E5	9	49
HPV31	E2	9	159		HPV31	E5	11	1
HPV31	E2	10	159		HPV31	E5	9	5
	E2	11	32		HPV31	E5	11	5 70
HPV31		8	158		HPV31	E5 E5	11	56
HPV31		10	158		HPV31		11	56
HPV31	E2	11	158					

Table X HLA A24 Supermotif Peptides

				TIEN NET Supermon	Lepudes			
HPV31	E5	9	31		HPV31	E6	11	69
HPV31		8	10		HPV31		10	95
HPV31		10	10		HPV31		9	61
HPV31		9	7		HPV31		10	61
HPV31		10	7		HPV31		8	118
			7		HPV31		11	118
HPV31		11			HPV31		9	11
HPV31		8.	35		HPV31	E6	11	11
HPV31		9	35					90
HPV31		10	35			E6	10	
HPV31		11	35		HPV31		11	90
HPV31		8	37		HPV31	E6	8	72
HPV31		9	37		HPV31		10	72
HPV31		10	37		HPV31		11	100
HPV31		8	41		HPV31		9	37
HPV31		9	41		HPV31		11	37
HPV31	E5	10	41		HPV31		9	91
HPV31	E5	8	8		HPV31		10	91
HPV31	E5	9	8		HPV31	E6	11	91
HPV31	E5	10	8		HPV31	E6	11	127
HPV31	E5	8	65		HPV31	E6	11	109
HPV31	E5	10	65		HPV31	E6	11	22
HPV31	E5	10	51		HPV31	E6	8	36
HPV31	E5	8	73		HPV31	E6	10	36
HPV31	E5	9	73		HPV31	E6	9	124
HPV31		11	47		HPV31	E6	9	68
HPV31		8	12		HPV31	E6	11	68
HPV31		11	12		HPV31	E6	11	27
HPV31		9	21		HPV31	E6	10	131
HPV31		10	33		HPV31		8	77
HPV31		11	33		HPV31		9	80
HPV31		8	64		HPV31		10	82
HPV31		9	64		HPV31		10	87
HPV31		11	64			E6	11	86
HPV31		8	38		HPV31		9	42
HPV31		9	38		HPV31		11	42
		11	38		HPV31		9	83
HPV31					HPV31		9	132
HPV31		8	50 50		HPV31		11	78
HPV31		11			HPV31		10	19
HPV31		9	63					
HPV31		10	63		HPV31		9	68 68
HPV31		9	46		HPV31		11	75
HPV31		9	18		HPV31 HPV31		9	75
HPV31		11	18					
HPV31		8	103		HPV31		10	75
HPV31		11	66		HPV31		8	21
HPV31		8	63		HPV31		9	14
HPV31		8	30		HPV31		10	48
HPV31		9	44		HPV31		9	81
HPV31	E6	11	44		HPV31		8	4
HPV31	E6	11	57		HPV31		10	4
HPV31	E6	10	75		HPV31		11	88
HPV31	E6	9	20		HPV31		10	78
HPV31	E6	8	25			E7	10	89
HPV31	E6	8	14		HPV31		8	82
HPV31	E6	10	14		HPV31	E7	8	6
HPV31		9	39		HPV31	E7	10	6
HPV31	E6	10	41		HPV31	E7	10	73
HPV31		8	47			E7	11	73
HPV31		8	69		HPV31	E7	8	77
HPV31		10	69		HPV31	E7	11	77

Table X HLA A24 Supermotif Peptides

					p	_		
HPV31	E7	11	66		HPV31	L1	8	141
HPV31		8	71		HPV31		10	141
HPV31		9	71		HPV31		11	266
HPV31		10	56		HPV31		9	115
HPV31		9	7		HPV31		10	115
HPV31		11	12		HPV31		8	474
HPV31		11	55		HPV31		8	307
HPV31		9	348		HPV31		9	376
HPV31		8	398		HPV31		9	399
HPV31		10	398		HPV31 HPV31		8	388
HPV31 HPV31		10 11	180 213		HPV31		10	382 382
HPV31		8	285		HPV31		11	382
HPV31		9	285		HPV31		9	181
HPV31		11	147		HPV31		11	181
HPV31		9	158		HPV31		9	61
HPV31		9	224		HPV31		11	61
HPV31		9	387		HPV31		10	33
HPV31		11	459		HPV31		11	33
HPV31		8	372		HPV31	L1	11	126
HPV31		11	372		HPV31		9	83
HPV31	L1	10	200		HPV31	L1	10	83
HPV31	L1	8	129		HPV31	L1	8	468
HPV31	L1	9	203		HPV31	L1	9	455
HPV31	L1	8	216		HPV31	L1	11	455
HPV31	L1	10	216		HPV31		8	381
HPV31	L1	9	353		HPV31		9	381
HPV31		8	336		HPV31		11	381
HPV31		11	417		HPV31		10	60
HPV31		9	151		HPV31		11	237
HPV31		10	151		HPV31		10	65
HPV31		10	234		HPV31		8	20
HPV31		9	444		HPV31 HPV31		8	231
HPV31 HPV31		8 10	370 370		HPV31		10 11	247 247
HPV31		8	107		HPV31		8	159
HPV31		8	449		HPV31		8	42
HPV31		9	363		HPV31		9	42
HPV31		11	363		HPV31		11	42
HPV31		8	26		HPV31		9	407
HPV31	L1	9	26		HPV31	L1	8	43
HPV31	L1	10	26		HPV31	L1	10	43
HPV31	L1	9	248	1	HPV31	L1	8	99
HPV31	Ll	10	248	1	HPV31	L1	10	3
HPV31		8	375		HPV31		11	3
HPV31		10	375		HPV31		11	389
HPV31		10	447		HPV31		10	238
HPV31		8	249		HPV31		11	238
HPV31		9	249		HPV31		9	201
HPV31		9	91		HPV31		11	201
HPV31		11	91		HPV31		8	300
HPV31		10	205		HPV31		9 11	300 32
HPV31		8	85		HPV31			
HPV31		8 9	323 323		HPV31		10 8	451 342
HPV31		8	117		HPV31			220
HPV31		11	117		HPV31		8	441
HPV31		10	105		HPV31			222
HPV31		11	68		HPV31		11	222
HPV31			406		HPV31		9 .	188
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Table X HLA A24 Supermotif Peptides

				TLEAT TEAT OUPCOMOUNT TO	pilaci	,		
HPV31	L1	8	464	HI	PV31	L1	11	446
HPV31	L1	11	113	HI	PV31	L1	8	28
HPV31	L1	11	17	H	2V31	L1	8	301
HPV31	L1	8	242	H	PV31	L1	10	334
HPV31	L1	9	242	H	V31	LI	8	62
HPV31	L1	10	242	H	2V31	L1	10	62
HPV31		9	374		V31		9	292
HPV31	L1	11	374	H	V31	L1	9	391
HPV31	L1	8	462		PV31			391
HPV31		9	452		PV31		11	277
HPV31		10	462		PV31		9	235
HPV31	L1	8	306		PV31		8	364
HPV31		9	306		V31		10	364
HPV31		11	378		V3 1		8	250
HPV31		11	156		V31		8	445
HPV31		8	255		V31		8	27
HPV31		9	70		V31		9	27
HPV31		8	420		V31		9	34
HPV31		9	41		V31		10	34
HPV31		10	41		V31		11	286
HPV31		8	77		V31		9	311
HPV31		10	77 98		V31		10 11	311
HPV31		9	75		V31		8	226
HPV31		10 10	75 90		V31		10	135
HPV31		8	228		V31		11	135
HPV31		9	228		V31		11	342
HPV31		11	228		V31		11	376
HPV31		11	51		V31		11	251
HPV31		8	414		V31		11	385
HPV31		11	2		V31		8	275
HPV31		9	149		V31		8	438
HPV31		11	149		V31		9	438
HPV31		10	394	HP	V31	L2	10	438
HPV31	L1	9	299		V31		11	438
HPV31		10	299	HP	V31	L2	10	278
HPV31	L1	10	283	HP	V31	L2	8	354
HPV31	L1	11	283	HP	V31	L2	8	116
HPV31	L1	11	23	HP	V31	L2	11	31
HPV31	L1	10	340		V31		8	190
HPV31	L1	11	290	HP	V31	L2	8	171
HPV31		8	194		V31		9	253
HPV31		8	271		V31		10	253
HPV31		10	355		V31		11	253
HPV31		11	355		V31		10	196
HPV31		8	286		V31		8	237
HPV31		11	246		V31		9	237
HPV31		9	383		V31			237
HPV31		10	383		V31		8	433
HPV31		8	354		V31			433
HPV31		11	354		V31			439
HPV31		8	408		V31 V31			439
HPV31		10	267		V31 V31			439
HPV31		11	267		V31 V31		11	113 351
HPV31		9	44		V31 V31		11 8	26
HPV31 HPV31		11	333 291		V31 V31			26
HPV31		10	390		V31			65
HPV31		11	390		V31			65
HPV31		10	418		V31		9	52
mPV31	TIT.	10	410	ne			-	-2

Table X HLA A24 Supermotif Peptides

				TILL TILL TO APPENDE	· r optide	•		
HPV31	L2	8	213		HPV31	L2	11	303
HPV31	L2	10	213		HPV31	L2	9	229
HPV31	L2	11	213		HPV31	L2	10	229
HPV31	L2	9	175		HPV31	L2	11	429
HPV31		8	38		HPV31		8	298
HPV31	L2	9	38		HPV31	L2	10	9
HPV31		11	38		HPV31		11	9 .
HPV31	L2	9	318		HPV31		8	306
HPV31		9	403		HPV31		8	316
HPV31		8	432		HPV31		11	316
HPV31		9	432		HPV31		8	47
HPV31		10	432		HPV31		9	47
HPV31		9	279		HPV31		11	295
HPV31		10	45		HPV31		8	94
HPV31		11	45		HPV31		9	431
HPV31		8	245		HPV31		10	431
HPV31		10	114		HPV31		11	431
HPV31		10	105		HPV31		9	325
HPV31		9	197		HPV31 HPV31		11	325 86
HPV31		11	35		HPV31	L2	10	182
HPV31		8	231		HPV31		11	104
HPV31 HPV31		9	176		HPV31		8	104
HPV31		10	287		HPV31		11	260
HPV31		10	352		HPV31		9	50
HPV31		10	261		HPV31		11	50
HPV31		11	447		HPV31		8	396
HPV31		8	269		HPV31		9	396
HPV31		9	269		HPV31		8	151
HPV31		11	269		HPV31		8	184
HPV31		8	204		HPV31		10	184
HPV31		10	390		HPV31	L2	8	346
HPV31		8	292		HPV31	L2	10	346
HPV31		10	187		HPV31	L2	11	346
HPV31	L2	11	187		HPV31	L2	8	379
HPV31		10	410		HPV31	L2	8	80
HPV31	L2	10	402		HPV31		10	80
HPV31	L2	11	210		HPV31	L2	10	149
HPV31	L2	11	122		HPV31		11	195
HPV31	L2	8	88		HPV31		9	236
HPV31	L2	11	422		HPV31		10	236
HPV31		9	100		HPV31		8	157
HPV31		8	394		HPV31	L2	9	40
HPV31		10	394		HPV31		8	312
HPV31		11	394		HPV31		9	312
HPV31		10	235		HPV31		10	312
HPV31		11	235		HPV31 HPV31		9 10	347
HPV31		9	156					304
HPV31		8	388		HPV31	L2	10 9	136
HPV31		10	167		HPV31		10	136
HPV31		11	167		HPV31		8	39
HPV31		9	415		HPV31	L2	10	39
HPV31		8	425		HPV31		8	426
HPV31		9	425 127		HPV31	L2	9	344
HPV31			97		HPV31	L2	10	344
HPV31 HPV31		9 10	97		HPV31	L2	10	343
HPV31		8	44		HPV31	L2	11	343
HPV31		11	44		HPV31		9	391
HPV31		10	243		HPV31		11	391
UPVJI	шZ	10	742					

Table X HLA A24 Supermotif Pentides

				HLA A24 Supe	rmotif Peptide	S		
HPV31	T.2	8	254		HPV33	R1	11	352
HPV31		9	254		HPV33		10	38
HPV31		10	254			E1	11	38
HPV31		8	392		HPV33	E1	11	295
HPV31		10	392		HPV33	E1	9	331
HPV31		9	81		HPV33	E1	9	605
HPV31		8	53		HPV33	E1	11	605
HPV31		10	32		HPV33	E1	10	50
HPV31		9	262		HPV33	E1	8	449
HPV31		8	440		HPV33	E1	9	449
HPV31	L2	9	440		HPV33	El	11	456
HPV31	L2	10	386		HPV33	E1	10	385
HPV31	L2	8	319		. HPV33	E1	8	212
HPV33	E1	9	452		HPV33	E1	8	446
HPV33	E1	9	448		HPV33	E1	11	446
HPV33	E1	10	448		HPV33	E1	11	501
HPV33	E1	11	384		HPV33	E1	8	265
HPV33		10	596		HPV33		10	265
HPV33	E1	11	596		HPV33		8	459
HPV33		9	532		HPV33		9	459
HPV33		10	546		HPV33		10	459
HPV33	E1	11	546		HPV33	E1	11	209
HPV33		9	311		HPV33		10	235
HPV33		9	81		HPV33	E1	8	11
HPV33		11	22		HPV33 HPV33		9 .	512 512
HPV33		9	207 259		HPV33		11	480
HPV33		8	259		HPV33		8	44
HPV33 HPV33		9 10	259		HPV33		8	564
HPV33		11	259		HPV33		9	564
HPV33		9	297		HPV33		10	327
HPV33		9	226		HPV33		9	16
HPV33		11	226		HPV33		11	256
HPV33		11	14		HPV33	E1	10	404
HPV33		8	118		HPV33	E1	10	347
HPV33	E1	11	118		HPV33	E1	9	266
HPV33	E1	8	494		HPV33	E1	8	267
HPV33	E1	9	494		HPV33	E1	9	200
HPV33	E1	10	494		HPV33		11	200
HPV33	E1	9	367		HPV33	E1	8	394
HPV33	E1	10	46			E1	10	435
HPV33		8	78		HPV33		8	203
HPV33		8	349		HPV33	E1	10	203
HPV33		10	349		HPV33		11	203
HPV33		8	62			E1	10	124
HPV33		9	62		HPV33		11	510
HPV33		11	62		HPV33		8	393 393
HPV33		10	541		HPV33 HPV33	E1	9	285
HPV33		9	324		HPV33	E1	9	304
HPV33		10	324		HPV33	E1	10	304
HPV33		9	516		HPV33		11	304
HPV33 HPV33		10 9	516 228		HPV33		8	412
HPV33		11	49		HPV33		8	425
HPV33		8	580		HPV33	E1	11	425
HPV33		9	445			E1	9	223
HPV33		11	537		HPV33		9	245
HPV33		8	361		HPV33		10	245
HPV33		10	361		HPV33		11	69
HPV33		11	361		HPV33	E1	8	247

Table X HLA A24 Supermotif Peptides

HPV33	E1	11	247	HP	V33	E1	8	549
HPV33	E1	10	483		V33		9	549
HPV33		11	271		V33		8	437
HPV33		9	47		V33		10	437
HPV33		9	438		V33		8	308
HPV33		9	290		V33		11	146
HPV33		11	290		V33		8	355
HPV33		8	286		V33		9	273
HPV33		8	260		V33		-	280 280
HPV33		9	260 260		V33 V33		11 11	575
HPV33		10 11	260		V33		9	335
HPV33 HPV33		9	362		V33		8	608
HPV33		10	362		V33		9	608
HPV33		11	362		V33		9	568
HPV33		10	281		V33		10	440
HPV33		11	281		V33		11	440
HPV33		10	576	HP	V33	E1	10	604
HPV33		8	336	HP	V33	E1	9	211
HPV33	E1	9	547	HP	V33	E1	9	342
HPV33		10	547	HP	V33	E1	9	539
HPV33	E1	11	547	HP	V33	E1	10	111
HPV33	E1	8	201		V33		9	292
HPV33	E1	10	201		V33			482
HPV33		8	253		V33		11	243
HPV33	E1	8	312		V33			252
HPV33		11	312 .		V33			239
HPV33		В	513		V33			239
HPV33		10	513		V33		8	521 521
HPV33		8	298		V33 V33		9	477
HPV33		10	353		V 3 3			477
HPV33		8	443		V 3 3		11	477
HPV33		11	346		V33		9	183
HPV33 HPV33		11	346		V33		9	328
HPV33		10	199	· HP			9	282
HPV33		9	71		V33		10	282
HPV33		10	289	HP*	V33	E1	9	577
HPV33		9	135	HP	V33	E1	11	577
HPV33		9	473	HP	V33	E1	11	337
HPV33	E1	11	473	HP4	V33	E1	8	609
HPV33	E1	8	195		V33		11	523
HPV33	E1	10	195		V33		10	119
HPV33		10	560		V33		8 .	578
HPV33		10	175		V33			578
HPV33		11	181		V33			23
HPV33		10	601		V33			491
HPV33		8	431		V33			190 246
HPV33		9	431		V33 V33			246
HPV33		11	431		V33 V33			338
HPV33 HPV33		10	586 519		V33			338
HPV33		11	519		V33			182
HPV33		8	434		V33			517
HPV33		11	434					517
HPV33		9	505			E1	9	594
HPV33		10	505			E1	8	17
HPV33		10	593			E1	9	314
HPV33		10	570	HP	V33	E1	8	332
HPV33		8	485	HP	V33	E1	10	502

Table X HLA A24 Supermotif Peptides

				THE STATE OF CHILDREN	сриче			
HPV33	E1	8	522	H	PV33	E2	8	35
HPV33	E1	9	478	н	IPV33	E2	9	35
HPV33		10	478	н	IPV33	E2	10	35
HPV33	E1	10	524	н	IPV33	E2	9	62
HPV33		9	571		PV33		8	42
HPV33		11	571		PV33		10	42
HPV33		11	528		PV33		11	240
HPV33		10	313		PV33		9	18
HPV33		9	69		PV33		9	299
HPV33		11	69	н	PV33	E2	11	299
HPV33		10	78		PV33		9	43
HPV33		9	41	н	PV33	E2	8	333
HPV33.		11	41		PV33		8	147
HPV33		9	10		IPV33		9	147
HPV33		10	10		PV33		11	147
HPV33		9	288		PV33		11	183
HPV33		10	145	н	PV33	E2	11	157
HPV33		11	145		PV33		8	127
HPV33		9	25		PV33		11	127
HPV33		11	25		PV33		8	272
HPV33		10	17		PV33		8	133
HPV33		9	235		PV33		9	196
HPV33		10	235		PV33		9	342
HPV33		9	232		PV33		10	342
HPV33		8	153		PV33		9	295
HPV33		8	130		PV33		8	29
HPV33		. 9	130		PV33		10	29
HPV33		10	130		PV33		9	345
HPV33		11	130		PV33		10	203
HPV33		9	74		PV33		9	332
HPV33		10	298	н	PV33	E2	8	96
HPV33		8	80		PV33		9	71
HPV33		9	185		PV33		9	191
HPV33		11	100		PV33		9	91
HPV33		9	325	н	PV33	E2	10	120
HPV33		10	325	н	PV33	E2	9	86
HPV33		11	325	н	PV33	E2	8	292
HPV33		11	336	н	PV33	E2	9	292
HPV33		10	53	н	PV33	E2	9	7
HPV33		11	53	н	PV33	E2	10	7
HPV33		9	278		PV33		8	37
HPV33		11	32	н	PV33	E2	11	266
HPV33		9	139	H	PV33	E2	10	290
HPV33	E2	11	276	н	PV33	E2	11	290
HPV33	E2	11	137	н	PV33	E2	9	285
HPV33	E2	8	339	H	PV33	E2	10	61
HPV33	E2	9	242	н	PV33	E2	8	302
HPV33	E2	9	34	н	PV33	E2	8 .	205
HPV33	E2	10	34	н	PV33	E2	10	324
HPV33	E2	11	34	H	PV33	E2	11	324
HPV33	E2	8	23	H	PV33	E2	10	93
HPV33	E2	11	23	H	PV33	E2	11	93
HPV33	E2	8	66	H	PV33	E2	10	128
HPV33		10	66	H	PV33	E2	11	128
HPV33		8	151	H	PV33	E2	9	146
HPV33	E2	9	151	H	PV33	E2	10	146
HPV33		10	151	H	PV33	E2	8	233
HPV33		10	169	H	PV33	E2	11	233
HPV33	E2	8	177		PV33		10	267
HPV33	E2	8	243	H	PV33	E2	10	337

Table X HLA A24 Supermotif Peptides

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HPV33	E2	8	343	HPV33	E5	9	5
HPV33	E2	9	343	HPV33	E5	11	5
HPV33	E2	11	343	HPV33	E5	8	49
HPV33	E2	8	72	HPV33	E5	9	49
HPV33	E2	11	72	HPV33	E5	10	49
HPV33	E2	8	192	HPV33	E5	11	49
HPV33	E2	8	326	HPV33	E5	8	2
HPV33	E2	9	326	HPV33	E5	9	2
HPV33	E2	10	326	HPV33	E5	10	2
HPV33	E2	11	323	HPV33	E5	11	2
HPV33	E2	8	148	HPV33	E5	8	10
HPV33		10	148	HPV33	E5	9	10
HPV33		11	148		E5	10	10
HPV33	E2	10	101	HPV33	E5	9	23
HPV33	E2	9	102	HPV33	E5	10	23
HPV33	E2	8	92	HPV33	E5	11	23
HPV33	E2	11	92	HPV33	E5	8	48
HPV33		9	170	HPV33	E5	9	48
HPV33	E2	9	159	HPV33	E5	10	48
HPV33	E2	10,	159	HPV33	E5	11	48
HPV33		10	138	HPV33	E5	8	11
HPV33	E2	8	44	HPV33	E5	9	11
HPV33	E2	11	44	HPV33	E5	8	55
HPV33	E2	В	131	HPV33	E5 E5	10 8	55 22
HPV33	E2	9	131 131	HPV33	E5	10	22
HPV33 HPV33	E2 E2	10 10	151	HPV33	E5	11	22
HPV33	E2	11	158	HPV33	E5	8	54
HPV33	E5	8	30	HPV33	E5	9	54
HPV33	E5	9	30	HPV33	E5	11	54
HPV33	E5	10	30	HPV33	E5	8	17
HPV33		11	30	HPV33	E5	10	17
HPV33		8	8	HPV33	E5	11	37
HPV33		10	8	HPV33	E5	9	18
HPV33		11	8	HPV33	E5	8	32
HPV33	E5	9	63	HPV33	E5	و	32
HPV33	E5	9	14	HPV33	E5	11	32
HPV33	E5	10	14	HPV33	E5	10	38
HPV33	E5	11	14	HPV33	E5	8	35
HPV33	E5	8	52	HPV33	E5	8	33
HPV33	E5	10	52	HPV33	E5	10	33
HPV33	E5	11	52	HPV33	E5	9	39
HPV33	E5	8	50	HPV33	E5	11	39
HPV33	E5	9	50	HPV33	E5	8	57
HPV33	E5	10	50	HPV33	E5	10	57
HPV33	E5	9	9	HPV33	E5	9	1
HPV33	E5	10	9	HPV33	E5	10	1
HPV33	E5	11	9	HPV33	E5	11	1
HPV33	E5	8	12	HPV33	E5	11	61
HPV33	E5	11	12	HPV33	E5	9	21
	E5	9	56	HPV33	E5	11	21
	E5	11	56	HPV33	E5	8	46
HPV33		8	3	HPV33	E5	9	46
HPV33		9	3	HPV33	E5	10	46
HPV33		10	3	HPV33	E5	11	46
	E5	11	3	HPV33		8	25
	E5	8	42	HPV33	E5 E5	9	25
HPV33	E5	9	42	HPV33 HPV33	E5	11	25
HPV33		10	42	HPV33		9	16 16
HPV33	RD	8	5	nevaa	EJ	9	10

Table X HLA A24 Supermotif Peptides

				TILA A2+ Supermoni	repudes	,		
HPV33	E5	11	16		HPV33	E6	10	36
HPV33	E5	9	27		HPV33	E6	10	90
HPV33		11	27		HPV33	E6	11 '	90
	E5	8	28		HPV33	E6	9	124
HPV33		10	28		HPV33	E6	9	68
HPV33		11	28		HPV33	E6	10	10
HPV33		9	41		HPV33	E6	10	131
HPV33		10	41		HPV33	E6	10	87
HPV33		11	41		HPV33	E6	9	11
HPV33		8	4		HPV33	E6	8	21
HPV33		9	4		HPV33	E6	9	91
HPV33		10	4		HPV33	E6	10	91
HPV33		8	6		HPV33	E6	11	91
HPV33		10	6		HPV33	E6	10	51
HPV33		9	34		HPV33		9	52
HPV33		8	31		HPV33		11	52
HPV33		9	31		HPV33	E6	9	42
HPV33		10	31		HPV33	E6	8	53
HPV33		8	40		HPV33	E6	10	53
HPV33		10	40		HPV33	E7	9	68
HPV33		11	40		HPV33	E7	8	75
HPV33		9	53		HPV33	E7	9	75
HPV33		10	53		HPV33	E7	10	75
HPV33		9	58		HPV33		8	21
HPV33		9	46		HPV33		9	14
HPV33		9	18		HPV33	E7	10	14
	E6	11	18		HPV33	E7	9	59
HPV33		8	103		HPV33	E7	8	82
HPV33		8	66		HPV33	E7	8	15
HPV33		11	66		HPV33	E7	9	15
HPV33		8	30		HPV33	E7	11	15
HPV33		11	139		HPV33	E7	10	19
HPV33		11	44		HPV33	E7	8	6
HPV33		10	14		HPV33	E7	10	6
HPV33		9	120		HPV33	E7	11	6
HPV33		9	4		HPV33		9	81
HPV33		8	98		HPV33		11	66
HPV33		11	27		HPV33	E7	8	77
HPV33		9	20		HPV33	E7	9	71
HPV33		10	41		HPV33	E7	9	7
HPV33		10	75		HPV33		10	7
HPV33		8	69		HPV33		11	12
HPV33		11	69		HPV33		10	290
HPV33		9	61		HPV33		10	392
HPV33		10	61		HPV33		9	44
HPV33		8	118		HPV33	L1	8	270
HPV33		11	118		HPV33		11	147
HPV33		11	78		HPV33		10	345
HPV33		8	72		HPV33	L1	9	223
HPV33		10	72		HPV33	L1	8	396
HPV33		10	64		HPV33		10	396
HPV33		11	100		HPV33		11	457
HPV33		11	50		HPV33		10	449
HPV33		11	86		HPV33		8	370
	E6	9	80		HPV33		10	199
HPV33		9	59		HPV33		8	129
HPV33		11	59		HPV33	L1	9	202
HPV33	E6	10	95		HPV33	L1	8	335
HPV33		11	95		HPV33	L1	10	335
HPV33		8	36		HPV33	L1	11	415
		-						

Table X HLA A24 Supermotif Peptides

				III.	······································			
HPV33	L1	9	151		HPV33	L1	8	306
HPV33		10	151		HPV33	L1	11	190
HPV33		10	233		. HbA33		8	42
HPV33		8	107		HPV33	L1	9	42
HPV33		8	447		HPV33		11	42
HPV33		9	385		HPV33		9	61
HPV33		á ·	368		HPV33		11	61
HPV33		10	368		HPV33		8	382
HPV33		9	361		HPV33		9	382
HPV33		11	361		HPV33		9	405
		8	26		HPV33		8	62
HPV33 HPV33		9	26		HPV33		10	62
		10	26		HPV33		8	99
HPV33 HPV33		11	26		HPV33		10	99
			247		HPV33		10	237
HPV33		9	247		HPV33		11	237
HPV33		10			HPV33		11	387
HPV33		8	248		HPV33		9	200
HPV33		9	248		HPV33		11	200
HPV33		8	260		HPV33		8	299
HPV33		9	291		HPV33		9	299
HPV33		8	249		HPV33		9	192
HPV33		10	373					
HPV33		10	445		HPV33		11	221
HPV33		9	91		HPV33		8	187
HPV33		11	91		HPV33		9	187
HPV33		10	204		HPV33		8	439
HPV33		8	85		HPV33		8	462
HPV33		9	322		HPV33		11	113
HPV33		8 1	117		HPV33		8	55
HPV33		11	117		HPV33		11	17
HPV33		10	105		HPV33		8	241
HPV33		8	472		HPV33		9	241
HPV33		11	68		HPV33		10	241
HPV33		10	404		HPV33		8	460
HPV33		11	265		HPV33		9	460
HPV33		11	281		HPV33		10	460
HPV33		9	115		HPV33		11	372
HPV33		10	115		HPV33		8	305
HPV33		9	259		HPV33		9	305
HPV33		9	365		HPV33		8	254
HPV33		11	365		HPV33		8	347
HPV33		9	397		HPV33		9	70
HPV33		10	33		HPV33		8	418
HPV33		11	33		HPV33		9	41
HPV33		11	126		HPV33		10	41
HPV33		9	83		HPV33		8	77
HPV33		10	83		HPV33		10	77
HPV33		8	466		HPV33		9	98
HPV33		9	453		HPV33		11	98
HPV33		11	453		HPV33		10	75
HPV33		10	60		HPV33		10	90
HPV33		11	236		HPV33		11	51
HPV33		10	65		HPV33		8	285
HPV33		9	379		HPV33		11	32
HPV33		11	379		HPV33		11	245
HPV33	L1	8	20		HPV33		8	412
HPV33	L1	8	230		HPV33		9	149
HPV33	L1	9	442		HPV33		11	149
HPV33	L1	10	246		HPV33		9	298
HPV33	L1	11	246		HPV33	61	10	298

Table X HLA A24 Supermotif Peptides

				HLA A24 Supermont P	epudes			
HPV33	L1	8	227	н	IPV33	L2	10	283
HPV33	Ll	9	227	H	IPV33	L2	10	272
HPV33	L1	11	227	H	IPV33	L2	11	272
HPV33	L1	11	23		IPV33		8	327
HPV33	L1	8	352	H	IPV33	L2	10	431
HPV33	L1	11	352		IPV33		10	264
HPV33		11	2		IPV33		10	401
HPV33		11	444		IPV33		9	350
HPV33		8	193		IPV33		10	95
HPV33		10	266		IPV33 IPV33		9 11	369 30
HPV33		11	266		IPV33		11	113
HPV33		11 9	212 381		(PV33		10	447
HPV33 HPV33		10	381		IPV33		8	242
HPV33		10	282		IPV33		9	242
HPV33		11	282		IPV33			242
HPV33		9	336		IPV33		8	440
HPV33		11	430	H	IPV33	L2	9	440
HPV33		11	332	H	IPV33	L2	10	440
HPV33		10	388	H	IPV33	L2	8	421
HPV33	L1	11	388		IPV33		10	421
HPV33		10	353		(PV33		8	25
HPV33		11	353		(PV33		9	75
HPV33		10	416		(PV33			75
HPV33		9	374		PV33		9	51
HPV33		8	386		(PV33		8 11	374 374
HPV33		8	380		PV33			336
HPV33		10 11	380 380		IPV33			336
HPV33		8	300		IPV33			323
HPV33		10	333		(PV33			284
HPV33		9	100		PV33			44
HPV33		10	3	H	[PV33	L2	11	44
HPV33	L1	11	3	H	(PV33	L2	9	448
HPV33	L1	9	389		IPV33		11	448
HPV33	L1	10	389		PV33		10	292
HPV33		11	276		PV33			250
HPV33		8	362		PV33			250 104
HPV33		10	362		(PV33	L2	10	104
HPV33		9	234 443		PV33		8	433
HPV33 HPV33		8	27		(PV33			433
HPV33		9	27			L2		248
HPV33		10	27			L2		311
HPV33		8	35	H	PV33	L2	11	34
HPV33		9	35	H	(PV33	L2	8	236
HPV33	L1	9	34	H	PV33	L2	8	46
HPV33	L1	10	34	H		L2		46
HPV33	L2	11	256			L2		414
HPV33		9	241			L2		414
HPV33		10	241				8	107
HPV33		11	291		PV33		9	249 243
HPV33		10	23		PV33	L2 L2		243
HPV33		11	308 385			L2		397
HPV33 HPV33		10 8	280		PV33		10	372
HPV33		8	439		PV33		10	391
HPV33		9	439			L2	10	143
HPV33		10	439			L2	8	209
HPV33		11	439	H	PV33	L2	8 .	426

Table X HLA A24 Supermotif Peptides

				TICK A24 Superinous I	cpade	,		
HPV33	L2	8	420	I	HPV33	L2	10	309
HPV33	L2	9	420	I	IPV33	L2	9	265
HPV33	L2	11	420	F	IPV33	L2	9	386
HPV33	L2	11	73		IPV33	L2	11	386
HPV33	L2	11	215		IPV33	L2	10	132
HPV33		8	423		IPV33		11	132
HPV33	L2	11	423		IPV33		8	93
HPV33	L2	11	333			L2	9	96
HPV33	L2	9	99		IPV33	L2	9	337
	L2	10	99		IPV33	L2	10	187
HPV33		9	413		IPV33		10	251
HPV33		10	347		IPV33	L2	8	52
HPV33		11	395		IPV33	L2	10	31
HPV33		9	376		IPV33 IPV33	L2 L2	9	441 441
HPV33 HPV33		10 9	79 161		IPV33		11	404
HPV33		9	416		IPV33	L2	11	131
HPV33		8	186		IPV33	L2	9	92
HPV33		11	186		(PV33	L2	9	434
HPV33		8	221		IPV33	L2	11	446
HPV33		8	403		IPV33	L2	8	324
HPV33		10	91		1PV33	L2	11	324
HPV33		8	317		IPV45	El	11	384
HPV33		9	317		IPV45	E1	9	532
HPV33		8	43	H	IPV45	E1	9	311
HPV33		11	43	H	IPV45	E1	9	199
HPV33	L2	11	153	H	IPV45	E1	10	199
HPV33	L2	9	234	H	IPV45	E1	11	512
HPV33	L2	10	234		IPV45	E1	8	40
HPV33	L2	11	300		IPV45	E1	11	40
HPV33	L2	8	321		IPV45	E1	8	517
HPV33		11	321		IPV45	E1	9	517
HPV33		9	388		IPV45	E1	10	251
HPV33		8	303		IPV45	E1	9	202
HPV33		8	259			E1	11	202
HPV33		10	259		PV45	E1 E1	10 9	604 259
HPV33		10	357		IPV45	E1	10	259
HPV33 HPV33		8	393 122		IPV45	E1	11	259
HPV33		9	151			E1	9	297
HPV33		11	103			E1	8	226
HPV33		9	49		PV45	El	9	226
HPV33		11	49			E1	10	634
HPV33		8	106			E1	8	521
HPV33		9	106	Н	PV45	E1	9	521
HPV33	L2	8	156	H	PV45	E1	9	49
HPV33	L2	8	38	H	(PV45	E1	10	206
HPV33	L2	10	38		PV45	E1	8	349
HPV33	L2	8	189		IPV45	E1	10	349
HPV33	L2	10	189	H	IPV45	E1	9	108
HPV33	L2	11	352			E1	11	108
HPV33		10	146			E1	8	361
HPV33	L2	10	192			E1	10	361
HPV33	L2	8	355			E1	9	367
HPV33	L2	9	355			E1	10	46
HPV33	L2	8	162			E1	11	352
HPV33	L2	9	39		PV45	E1 E1	8	106 106
HPV33	L2	10	154			E1	11	623
HPV33	L2	9	432			E1	9	42
HPV33	L2	11	432	н	rv45	ET.	9	*4

Table X HLA A24 Supermotif Peptides

				•				
HPV45		10	42		HPV45	E1	9	292
HPV45	E1	9	328		HPV45		10	404
HPV45		11	328		HPV45		11	338
HPV45		8	431		HPV45		11	491
HPV45		9	431		HPV45		8	184
HPV45		11	431		HPV45		10	23
HPV45		9	445		HPV45		9	207
HPV45		10	596		HPV45		8	425
HPV45		11	596		HPV45	E1	11	425
HPV45		10	38		HPV45	E1	9	304
HPV45	E1	11	295		HPV45	E1	10	304
HPV45		9	74		HPV45	E1	11	304
HPV45	E1	11	74		HPV45	E1	9 .	245
HPV45	E1	9	324		HPV45	E1	10	245
HPV45	E1	10	324		HPV45	E1	9	223
HPV45	E1.	8	331		HPV45	E1	11	223
HPV45		9	331		HPV45 HPV45	E1 E1	9	263
HPV45	E1 E1	11	605		HPV45	E1	10 9	263
HPV45	E1	3.1	605 50		HPV45	E1	9	393 129
HPV45	E1	9	483		HPV45	E1	8	247
HPV45	E1	10	483		HPV45	E1	11	247
HPV45	E1	8	446		HPV45	E1	8	267
HPV45	E1	11	446		HPV45	E1	9	290
HPV45	E1	11	456		HPV45	E1	10	290
HPV45	E1	10	385		HPV45	E1	11	290
HPV45	E1	11	385		HPV45	E1	11	190
HPV45	E1	9	486		HPV45	E1	9	547
HPV45	E1	8	449		HPV45	E1	10	547
HPV45	E1	9	449		HPV45	E1	11	547
HPV45	E1	9	438		HPV45	E1	11	271
HPV45	E1	10	438		HPV45	E1	9	362
HPV45	E1	8	212		HPV45	E1	11	362
HPV45	E1	9	579		HPV45	E1	8	336
HPV45	E1	8	130		HPV45	E1	10	281
HPV45	E1	8	494		HPV45	E1	11	281
HPV45	E1 .	9	494		HPV45	E1	8	253
HPV45	E1	10	494		HPV45	E1	11	468
HPV45	E1	11	243		HPV45	E1	8	312
HPV45	E1	9	342		HPV45	E1	11	312
HPV45	E1	11	209		HPV45	E1	8	200
HPV45	E1	8	480		HPV45	E1	9	200
HPV45	E1	8	11		HPV45	E1	11	200
HPV45	E1	9	11		HPV45	E1	10	513
HPV45	E1	8	459		HPV45	E1	8	298
HPV45	E1	9	459		HPV45	E1	9	47
HPV45	B1	10	459		HPV45	E1	11	47
HPV45	E1	11	459		HPV45	E1	10	353
HPV45	E1	8	443		HPV45	E1	10	347
HPV45	E1	11	443		HPV45	E1	10	560
HPV45	E1	8	265		HPV45	E1	8	414
HPV45	E1	10	265		HPV45	E1	9	473
HPV45		10	235		HPV45	E1	11	473
HPV45	E1	8	16		HPV45	E1	8	177
HPV45	E1 E1	8 10	485 485		HPV45	E1 E1	11 10	177 601
					HPV45	E1	10	586
HPV45 HPV45	E1	9	256		HPV45	E1	8	554
	E1 E1	10 11	519 519		HPV45	E1	11	537
HPV45	E1	8	292		HPV45	E1	8	434
nPV45	E.I	0	232		11FV43	m T	u	434

Table X HLA A24 Supermotif Peptides

			TIER RE4 Superino	in replaces		
HPV45	E1.	9	505	HPV45 E1	9	577
HPV45	E1	10	505	HPV45 E1	11	577
HPV45	E1	10	546	HPV45 E1	11	523
HPV45	E1	11	546	HPV45 E1	8	203
HPV45	E1	10	238	HPV45 E1	10	203
HPV45	E1	11	238	HPV45 E1	8	578
HPV45	E1	10	593	HPV45 E1	10	578
	E1	10	570	HPV45 E1	9	81
HPV45	E1	10	437	HPV45 E1	9	266
HPV45	E1	11	437	HPV45 E1	9	635
	El	8	549	HPV45 E1	10	576
HPV45	E1	9	549	HPV45 E1	9	594
HPV45	E1	8	102 .	HPV45 E1	9	418
HPV45	E1	8	412	HPV45 E1	8	332
HPV45	E1	10	412	HPV45 E1	8	522
HPV45	E1	10	80	HPV45 E1	9	314
HPV45	E1	8	355	HPV45 E1	10	524
HPV45	E1	10	417	HPV45 E1	9	478.
HPV45	E1	10	128	HPV45 E1	10	478
	E1	9	335	HPV45 E1	9	571
HPV45	E1	8	280	HPV45 E1	11	571
HPV45	E1	11	280	HPV45 E1	8	394
HPV45	E1	8	608	HPV45 E1	11	528
HPV45	E1	11	575	HPV45 E1	10	313
	E1	9	273	HPV45 E2	8	78
	E1	8	440	HPV45 E2	11	78
HPV45	E1	10	440	HPV45 E2	11	47
HPV45	E1	11	440	HPV45 E2	10	84
HPV45	E1	10	482	HPV45 E2	10	16
HPV45	E1	11	482	HPV45 E2	10	247
HPV45	E1	9	448	HPV45 E2	8	216
HPV45	E1	10	448	HPV45 E2	9	305
HPV45	El	9	211	HPV45 E2	10	134
HPV45	E1	9	493	HPV45 E2	11	134
HPV45	E1	10	493	HPV45 E2	8	158
HPV45	E1	11	493	HPV45 E2	9 .	158
HPV45	E1	9	539	HPV45 E2	10	158
	E1	10	539	HPV45 E2	8	31
HPV45		8	183	HPV45 E2	9	31
HPV45	E1	9	183	HPV45 E2	11	31
	E1	11	288	HPV45 E2	8	171
HPV45	E1	8	308	HPV45 E2	8	212
HPV45	E1	10	104	HPV45 E2	11	351
	E1	8	477	HPV45 E2	8	319
	E1	10	477	HPV45 E2	9	80
HPV45		11	477	HPV45 E2	11	106
HPV45	E1	8	580	HPV45 E2	8	343
HPV45	E1	11	22	HPV45 E2	9	192
HPV45	E1	8	246	HPV45 E2	11	97
HPV45	E1	9	246	HPV45 E2	8	50
HPV45	E1	10	289	HPV45 E2	11	50
HPV45	E1	11	289	HPV45 E2	9	295
HPV45	E1	9	252	HPV45 E2	11	325
HPV45	E1	8	224	HPV45 E2	9	24
HPV45	E1	10	224	HPV45 E2	9	316
HPV45	E1	11	224	HPV45 E2	11	316
HPV45	E1	9	239	HPV45 E2	11	293
HPV45	E1	10	239	HPV45 E2	10	48
HPV45	E1	9	282	HPV45 E2	11	151
HPV45	E1	10	282	HPV45 E2	9	143
	-	-				

Table X HLA A24 Supermotif Peptides

HPV45	E2	10	143	HPV45	E2	8	199
	E2	10	59	HPV45	E2	9	353
HPV45	E2 :	9	2	HPV45	E2	10	352
HPV45		8	154	HPV45	E2	8	138
HPV45	E2	9	154	HPV45	E2	9	138
HPV45	E2	10	284	HPV45	E2	8	139
HPV45	E2	11	284	HPV45	E2	11	125
HPV45		9	312	HPV45	E2	11	164
HPV45	E2 -	8	184	HPV45		10	326
HPV45	E2	9	92		E2	11	326
HPV45	E2	9	49	HPV45	E2	10	98
HPV45		8	41	HPV45	E2	11	98
HPV45		9	41 .	HPV45	E2	10 9	126
HPV45		10	41	HPV45 HPV45	E2 E2	-	166 166
HPV45		10	107	HPV45	E2	10	145
HPV45		11	69	HPV45	E2	11	175
HPV45		9	109	HPV45	E2	8	137
HPV45			347	HPV45	E2	9	137
HPV45		9	332 265	HPV45	E2	10	137
HPV45		9	265	HPV45	E2	11	38
HPV45		8	289	HPV45	E2	10	165
HPV45		9	198	HPV45	E2	11	165
HPV45		8	136	HPV45	E2	8	144
HPV45	E2	9	136	HPV45	E2	9	144
HPV45		10	136	HPV45	E6	9	48
HPV45		11	136	HPV45	E6	9	37
HPV45		9	177	HPV45	E6	11	37
HPV45		9	360	HPV45	E6	9	61
HPV45		8	35	HPV45	E6	11	61
HPV45		10	35	HPV45	E6	11	59
HPV45		9	40	HPV45	E6	8	68
HPV45		10	40	HPV45	E6	11	68
HPV45	E2	11	40	HPV45	E6	8	105
HPV45		11	4	HPV45	E6	8	18
HPV45	E2	8	63	HPV45	E6	8	32
HPV45	E2	8	43	HPV45	E6	9	70
HPV45	E2	9	309	HPV45	E6	11	70
HPV45	E2	9	13	HPV45	E6	10	16
HPV45	E2	10	13	HPV45	E6	10	51
HPV45	E2	10	263	HPV45	E6	8	27
HPV45	E2	11	263	HPV45	E6	11	20
HPV45	E2	11	307	HPV45	E6	10	77
HPV45		10	142	HPV45	E6	10	97
HPV45		11	142	HPV45	E6	11	88
HPV45		9	302	HPV45	E6	10	43
HPV45		9	9	HPV45	E6	11	43
HPV45	E2	9	235	HPV45	E6	8	53
HPV45	E2	8	358	HPV45	E6	10	53
HPV45		9	358	HPV45	E6	8	71
	E2	11	358	HPV45	E6	10	71
	E2	8	160	HPV45	E6	11	71
HPV45		8	37	HPV45	E6	8	120
HPV45		8	348	HPV45	E6	11	120
HPV45		8	354	HPV45	E6	9	93 93
HPV45		9	99	HPV45	E6 E6	10 11	93
	E2	10	99	HPV45 HPV45	E6	9	93 54
HPV45	E2	11	213	HPV45	E6	8	92
HPV45	E2	11	190	HPV45	E6	10	92
HPV45	E2	11	339	UEA42	20	10	92

Table X HLA A24 Supermotif Peptides

				 ,	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,			
HPV45	E6	11	92	HP	V45	L1	8	103
HPV45	E6	9	13		V45		10	103
HPV45	E6	9	52		V45		11	28
HPV45	E6	11	52		V45		8	88
HPV45	E6	11	102	HP	V45	L1	10	88
HPV45	E6	9	98		V45		11	94
HPV45	E6	8	95		V4 5		9	184
HPV45	E6	9	95		V45			276
HPV45	E6	9	22		V45			409
HPV45		11	111		V45			188
HPV45		8	7		V45			318
HPV45		11	7		V45			250
HPV45		11	11		V45			488
HPV45		8	46		V45			480
HPV45		9	4.5		V45			401
HPV45		11	46		V45			401
HPV45		10	3		V45			301
HPV45		9	126		V45			226 _. 229
HPV45		8	74		V45 V45			242
HPV45 HPV45		9	41 79		V45			242
HPV45		11	29		V45			461
HPV45		11	24		V45			364
HPV45		10	84		V45			364
HPV45		10	89		V45			296
HPV45		11	89		V45			296
HPV45		8	38		745			169
HPV45		10	38	HP	V45	L1	9	177
HPV45	E6	9	85	HP	V45	L1	10	177 -
HPV45	E6	9	44	HP.	V45	L1	10	260
HPV45	E6	10	44	HP'	V45	L1	8	52
HPV45	E6	11	44	HP	V45			52
HPV45	E6	8	55		V45			52
HPV45		11	80		V45			133
HPV45		8	6		V45			133
HPV45		10	6		745			313
HPV45		10	64		V45			416
HPV45		8	25		V45			246
HPV45		8	83		745 745			399 399
HPV45 HPV45		9 10	83 83		745			274
HPV45		8	20		V45			274
HPV45		10	20		V45			404
HPV45		11	74		745			404
HPV45		11	91		745			14
HPV45		8	14	HP	745	L1	10	14
HPV45		11	11		745			14
HPV45		8	90	HP	V45	L1	8	287
HPV45	E7	10	75	HP	V45	L1	10	476
HPV45	E7	10	23	HP	745	L1	9	60
HPV45	E7	9	89	HP	V45	L1		60
HPV45	E7	8	85	HP	J45			351
HPV45	E7	11	85		J 45			351
HPV45		9	93		745			141
HPV45		9	7		V45			141
HPV45		10	86		V45			111
HPV45		8	94		V45			503
HPV45		9	76		V45			143
HPV45		10	12		745 745			143 131
HPV45	Ll	11	191	HP	v+5	PT	10	131

Table X HLA A24 Supermotif Peptides

				 	-		
HPV45	L1	9	199	HPV45	L1	11	4
HPV45	Ll	11	292	HPV45	L1	11	310
HPV45	Ll	10	435	HPV45	L1	11	49
HPV45	L1	10	231	HPV45	L1	9	219
HPV45	L1	9	286	HPV45	L1	8	383
HPV45	L1	9	396	HPV45	L1	11	383
HPV45	Ll	10	396	HPV45	L1	9	19
HPV45	L1	11	396	HPV45	L1	11	19
HPV45	Ll	9	392	HPV45	L1	8	17
HPV45	L1	11	392	HPV45	L1	9	17
HPV45	L1	9	13	HPV45	L1	11	17
HPV45	L1	11	13	HPV45		11	173
HPV45		8	23	HPV45	L1	8	22
HPV45		9	405	HPV45	L1	9	22
HPV45		10	59	HPV45	L1	8	248
HPV45		11	59	HPV45	L1	11	248
HPV45		10	12	HPV45	L1	8	214
HPV45		11	11 .	HPV45	L1	9	214
	L1	8	5	HPV45	L1	11	139
HPV45		9	5	HPV45	L1	9	440
HPV45		10	5	HPV45		11	440
HPV45		11	5	HPV45	Ll	11	380
HPV45		8	185	HPV45	L1	8	470 268
HPV45		8	411	HPV45	L1 L1	8 10	268
HPV45 HPV45		10 11	411	HPV45	L1	9	403
HPV45		8	328	HPV45	L1	11	403
	L1	9	328	HPV45		11	182
HPV45		8	6	HPV45		8	281
HPV45		9	6	HPV45		8	334
HPV45		10	6	HPV45	Ll	9	334
HPV45		9	117	HPV45	L1	10	206
HPV45	L1	11	117	HPV45	L1	11	263
HPV45	L1	9	109	HPV45	L1	8	491
HPV45	L1	10	109	HPV45	L1	9	491
HPV45	L1 -	8	497	HPV45	L1	9	96
HPV45	L1	9	484	HPV45	L1	9	67
HPV45	L1	11	484	HPV45	L1	10	67
HPV45		11	475	HPV45		9	124
HPV45		9	473	HPV45	L1	11	124
HPV45		10	91	HPV45	L1	10	101
HPV45		8	257		L1	8	46
HPV45		8	335	HPV45	L1	9	254 254
HPV45		8	68	HPV45 HPV45	L1 L1	11	254
HPV45 HPV45		9	68 413		L1	11	58
HPV45		9	413	HPV45	L1	8	427
HPV45		8	69		L1	9	327
HPV45		8	125		Li	10	327
HPV45		10	125	HPV45	L1	8	443
HPV45		10	29		ь1	11	272
HPV45		11	29	HPV45		10	423
HPV45		10	302	HPV45		11	115
HPV45		10	273		L1	10	376
HPV45		11	273		L1	11	43
HPV45	L1	9	227	HPV45	L1	9	175
HPV45	L1	11	227	HPV45	L1	11	175
HPV45	L1	8	4	HPV45	L1	10	419
HPV45		9	4		L1	11	419
HPV45	L1	10	4	HPV45	ы	8	220

Table X HLA A24 Supermotif Peptides

				Tan Time Captinous Topicaes		
	L1	9	410	HPV45 L2	8	319
HPV45	L1	11	410	HPV45 L2	9	51
HPV45	L1	10	116	HPV45 L2	8	430
HPV45	Ll	8	200	HPV45 L2	9	430
HPV45	L1	11	239	HPV45 L2	10	430
HPV45	L1	9	412	HPV45 L2	8	25
HPV45	L1	10	412	HPV45 L2	10	206
HPV45	L1	10	462	HPV45 L2	10	183
HPV45	Ll	9	365	HPV45 L2 HPV45 L2	9	433
HPV45	L1 L1	8	329 441	HPV45 L2 HPV45 L2	10 11	433 433
HPV45	L1	8	441	HPV45 L2	8	37
HPV45	L1	10 8	478	HPV45 L2	9	37
HPV45	L1	8	297	HPV45 L2	11	37
HPV45	Li	11	361	HPV45 L2	8	134
HPV45	Li	10	384	HPV45 L2	10	134
HPV45		8	20	HPV45 L2	8	292
HPV45	Li	10	20.	HPV45 L2	10	191
HPV45	Li	11	20	HPV45 L2	9	318
HPV45	L1	10	293	HPV45 L2	8	52
HPV45	Ll	11	293	HPV45 L2	9	406
HPV45	L1	8	417	HPV45 L2	10	406
HPV45	L1	10	362	HPV45 L2	9	279
HPV45	L1	9	126	HPV45 L2	8	407
HPV45	L1	10	319	HPV45 L2	9	407
HPV45	L1	9	477	HPV45 L2	9	44
HPV45	L1	9	420	HPV45 L2	10	44
HPV45	Ll	10	420	HPV45 L2	11	44
HPV45	L1	9	303	· HPV45 L2	10	143
HPV45	L1	9	261	HPV45 L2	8	400
HPV45	L1	8	53	HPV45 L2	9	400
HPV45	L1	9	53	HPV45 L2	8	43
HPV45	L2	8	161	HPV45 L2	10	43
HPV45	L2	11	286	HPV45 L2	11	43
HPV45	L2	10	328	HPV45 L2 HPV45 L2	11 8	34 346
HPV45 HPV45	L2	11	328 303	HPV45 L2 HPV45 L2	10	337
HPV45	L2 L2	11 8	340	HPV45 L2 HPV45 L2	11	337
HPV45	L2	11	340	HPV45 L2	10	242
	L2	9	255	HPV45 L2	11	375
	L2	11	255	HPV45 L2	9	392
HPV45	L2	8	275	HPV45 L2	10	392
HPV45	L2	10	405	HPV45 L2	9	248
	L2	11	405	HPV45 L2	8	438
	L2	10	278	HPV45 L2	9	305
HPV45	L2	11	322	HPV45 L2	8	270
HPV45	L2	11	142	HPV45 L2	10	270
HPV45	L2	9	345	HPV45 L2	11	270
HPV45	L2	11	83	HPV45 L2	10	387
HPV45	L2	11	30	HPV45 L2	8	325
HPV45	L2	10	397	HPV45 L2	10	325
	L2	11	397	HPV45 L2	8	399
	L2	8	331	HPV45 L2	9	399
	L2	8	241	HPV45 L2	10	399
	L2	11	241	HPV45 L2	8	258
	L2	9	122	HPV45 L2	11	336
	L2	11	157	HPV45 L2	10	391
HPV45	L2	8	306	HPV45 L2	11	391
	L2	8	368	HPV45 L2	9	98
HPV45	L2	10	368	HPV45 L2	9	120

Table X HLA A24 Supermotif Peptides

				nLA A24 Supermon	replice	•		
HPV45	T.2	11	120		HPV45	L2	10	217
HPV45		9	420		HPV45		11	217
HPV45		10	420		HPV45		10	427
HPV45		8	86		HPV45		11	427
HPV45		8	185		HPV45		9	369
HPV45		11	185		HPV45		10	31
HPV45		10	267		HPV45		8	249
HPV45		11	267		HPV45		11	249
HPV45		8	145		HPV45		9	388
HPV45		11	216		HPV45		11	112
HPV45		9	95		HPV45		9	444
HPV45		11	118		HPV45		11	444
HPV45		8	453		HPV45		9	428
HPV45		9	240		HPV45		10	428
HPV45		8	312		HPV45		11	428
HPV45		9	312		HPV45		8	437
HPV45		9	172		HPV45		9	437
HPV45		10	172		HPV45		8	401
HPV45		9	233		HPV45		10	443
HPV45		10	233		HPV45		8	436
HPV45		8	46		HPV45		9	436
HPV45		9	46		HPV45		10	436
HPV45		8	435		HPV45		11	442
HPV45		9	435		HPV56		8	15
HPV45		10	435		HPV56		11	15
		11	435		HPV56		9	21
HPV45					HPV56		10	21
HPV45 HPV45		11 10	295 451		HPV56		9	52
HPV45			298		HPV56		9	71
		9	298		HPV56		10	71
HPV45		8	316		HPV56		11	71
HPV45		11	316		HPV56		8	204
		8	220		HPV56		9	113
HPV45		8	235		HPV56		8	39
HPV45		9	367		HPV56		8	263
HPV45		11	367		HPV56		11	43
HPV45		9	49		HPV56		8	288
HPV45		11	49		HPV56		9	128
HPV45		10	247		HPV56		10	128
HPV45		9	362		HPV56		9	17
HPV45		9	154		HPV56		11	34
HPV45		11	358		HPV56		11	294
HPV45		8	424		HPV56		8	261
HPV45		9	149		HPV56		10	261
HPV45		9	384		HPV56		8	94
HPV45		10	250		HPV56		9	94
HPV45		8	121		HPV56		11	87
HPV45		10	121		HPV56		9	239
HPV45		8	363		HPV56		8	130
HPV45		11	363		HPV56		8	297
HPV45		9	39		HPV56		10	297
HPV45		10	304		HPV56		10	269
HPV45		10	376		HPV56		11	126
HPV45		8	38		HPV56		11	284
HPV45		10	38		HPV56		9	29
		8	136		HPV56		9	80
HPV45			359		HPV56		11	100
HPV45		10 9	135		HPV56		8	120
HPV45		11	426		HPV56		8	299
HPV45		8	389		HPV56		10	299
HPV45	LZ	0	202		112 7 3 0			

Table X HLA A24 Supermotif Peptides

				TIOITIE - Owponinous	r opiide.			
HPV56	E2	11	258		HPV56	E6	8	50
HPV56	E2	8	233		HPV56	E6	8	33
HPV56	E2	8	90		HPV56	E6	8	106
HPV56	E2	11	90		HPV56	E6	11	60
HPV56	E2	11	78		HPV56	E6	8	28
HPV56	E2	9	260		HPV56	E6	9	83
HPV56	E2	11	260		HPV56	E6	9	23
HPV56	E2	8	46		HPV56	E6	10	52
HPV56	E2	10	44		HPV56	E6	8	39
HPV56	E2	9	216		HPV56	E6	10	39
HPV56	E2	11	149		HPV56	E6	8	20
HPV56	E2	8	277		HPV56	E6	10	20
HPV56	E2	9	277		HPV56	E6	10	44
HPV56	E2	8	152		HPV56	E6	8	72
HPV56	E2	8	301		HPV56	E6	10	72
HPV56	E2	11	19		HPV56	E6	11	72
HPV56	E2	11	6		HPV56	E6	10	134
HPV56	E2	8	73 -		HPV56	E6	8	17
HPV56	E2	9	73		HPV56		10	17
HPV56	E2	10	73		HPV56	E6	11	17
HPV56	E2	8	253		HPV56	E6	9	94
HPV56	E2	9	253		HPV56	E6	10	94
HPV56	E2	9	246		HPV56	E6	11	94
HPV56	E2	10	251		HPV56	E6	8	54
HPV56	E2	11	251		HPV56		10	54
HPV56	E2	8	293		HPV56	E6	8	75
HPV56		9	272		HPV56	E6	10	75
	E2	10	272			E6	10	78
HPV56		10	26		HPV56		9	71
HPV56		8	141		HPV56	E6	11	71
HPV56		8	28		HPV56	E6	11	130
HPV56		10	28		HPV56	E6 E6	10 11	26 103
	E2	10	259		HPV56		10	70
HPV56		9	36		HPV56 HPV56	E6 E6	8	113
HPV56		10	36		HPV56	E6	9	40
HPV56		11 9	36 27		HPV56	E6	9	55
HPV56		11	27		HPV56	E6	9	47
HPV56		11	237		HPV56		11	47
HPV56		11	62		HPV56	E6	11	25
HPV56		10	63		HPV56	E6	9	112
HPV56		9	45		HPV56	E6	9	62
HPV56		10	35		HPV56	E6	10	62
HPV56		11	35		HPV56	E6	11	62
HPV56		9	270		HPV56	E6	10	98
HPV56		11	270		HPV56	E6	10	119
HPV56		10	79		HPV56	E6	9	127
HPV56		11	111		HPV56	E6	11	30
HPV56		8	74		HPV56	E6	8	80
HPV56		9	74		HPV56	E6	10	93
HPV56		9	102		HPV56	E6	11	93
HPV56		10	102		HPV56	E6	9	14
HPV56		8	81		HPV56	E6	11	14
HPV56		10	101		HPV56	E6	10	85
HPV56		11	101		HPV56	E6	10	90
HPV56	E6	11	89		HPV56	E6	9	21
HPV56		8	64		HPV56	E6	11	21
	E6	9	64		HPV56	E6	9	86
HPV56	E6	10	64		HPV56	E6	9	45
HPV56	E6	11	69		HPV56	E6	11	45

Table X HLA A24 Supermotif Peptides

HPV56	E6	8	56	F	IPV56	L1	9	419
HPV56	E6	9	135	F	IPV56	L1	8	402
HPV56	E6	11	81	F	IPV56	Ll	10	402
HPV56	E7	8	93	E	IPV56	L1	10	21
HPV56	E7	9	75	E	IPV56	L1	8	407
HPV56	E7	8	22	E	IPV56	L1	10	407
HPV56	E7	8	82	F	IPV56	L1	10	479
HPV56	E7	9	82	F	IPV56	L1	9	69
HPV56	E7	10	82	E	IPV56	L1	10	69
HPV56	E7	10	20	F	IPV56	L1	8	282
HPV56	E7	8	14	F	IPV56	L1	9	282
HPV56	E7	10	14	H	IPV56	L1	10	238
HPV56	E7	9	62	F	IPV56	L1	8	356
HPV56	E7	8	69	H	IPV56	L1	9	356
HPV56	E7	10	4	F	IPV56	L1	10	138
HPV56	E7	11	60	F	IPV56	L1	8	118
HPV56	E7	11	90	H	IPV56	L1	8	150
HPV56	E7	9	15 -	H	IPV56	L1	11	150
HPV56	E7	8	6	H	IPV56	L1	10	438
HPV56	E7	10	6	H	IPV56	L1	9	399
HPV56	E7	11	73	- H	IPV56	L1	11	399
HPV56	E7	8	84	H	IPV56	L1	8	15
HPV56	E7	11	84	H	IPV56	L1	11	20
HPV56	E7	9	7	H	IPV56	L1	8	32
HPV56	E7	10	12	H	IPV56	L1	10	68
HPV56	E7	11	11	H	IPV56	L1	11	68
HPV56	E7	10	85	H	IPV56	L1	8	414
HPV56	L1	11	58	H	IPV56	L1	10	414
HPV56	L1	8	381	H	IPV56	Ll	11	414
HPV56	L1	8	327	H	IPV56	L1	8	334
HPV56	L1	8	444		IPV56	L1	8	192
HPV56	L1	10	444		IPV56	L1	8	258
HPV56	L1	11	37		IPV56	L1	11	258
HPV56	L1	8	26		IPV56	L1	9	124
HPV56	L1	9	26			L1	11	124
HPV56	L1	11	26		IPV56	L1	9	8
HPV56		8 .	275		[PV56	L1	11	8
HPV56		9	275			L1	9	116
HPV56		10	275			L1	10	116
HPV56		10	422		IPV56		11	478
HPV56		11	422			L1	9	413
HPV56	L1	11	101			L1	11	413
HPV56		9	191			L1	11	270
	L1	9	195			L1	10	93
	L1	9	257		PV56	L1	10	300
HPV56	ь1	11	491			L1	11	300
HPV56	L1	10	233			L1	10	9 8 55
HPV56	L1	9	236			L1	8	
HPV56	L1	8	369			L1 L1	10	387 387
	L1	10	369				11 9	476
HPV56	L1	8	23			L1		264
HPV56	L1	10	23			L1 L1	8 10	450
	L1	11	23			L1	8	340
	L1	8	481			L1	11	224
	L1	10	267			L1	8	77
HPV56	L1	8	404			LI	9	77
HPV56	L1	11	404			L1	9	431
HPV56	Ll	8	303			L1	8	132
HPV56	L1	9	303			Ll	9	234
HPV56	LI	8	140	н	E V 30	DI	-	~34

Table X HLA A24 Supermotif Peptides

				TILA A24 Supermout	repude	S		
HPV56	L1	11	234		HPV56	L1	10	110
HPV56		8	333		HPV56	L1	9	131
HPV56		9	333		HPV56	L1	10	108
HPV56	L1	8	2		HPV56	L1	8	452
HPV56	Ll	8	1		HPV56	L1	9	487
HPV56	L1	9	1		HPV56	L1	11	487
HPV56	L1	10	5		HPV56	L1	11	67
HPV56	L1	10	376		HPV56	L1	8	446
HPV56	Ll	10	280		HPV56	L1	9	332
HPV56	Ll	11	280		HPV56	L1	10	332
HPV56	L1	9	6		HPV56		11	279
HPV56	L1	11	6		HPV56	L1	10	379
HPV56	L1	8	95		HPV56	L1	8 .	261
HPV56		10	95		HPV56		9	261
HPV56	L1	11	180		HPV56		11	261
HPV56	L1	10	123		HPV56	L1	9	489
HPV56		8	167		HPV56		9	182
HPV56	L1	8	430		HPV56		11	182
HPV56	L1	10	430		HPV56		11	86
HPV56		10	483		HPV56		11	323
HPV56		10	426		HPV56		8	61
HPV56		11	375		HPV56		9	61
HPV56		9	226		HPV56		10	61
HPV56		10	226		HPV56		8	304
HPV56		9	28		HPV56		9	377
HPV56		10	172		HPV56		9	415
HPV56		8	228		HPV56		10	415
HPV56		9	31		HPV56		8	215
HPV56		8	473		HPV56		10	215
HPV56		9	221		HPV56		11	215
HPV56		11	255		HPV56		9	370
HPV56		11	146		HPV56		11	366
HPV56		8	496		HPV56		10	38
HPV56		10	496		HPV56		11	38
HPV56		8	13		HPV56		8	29
HPV56		9	13		HPV56		11	29
HPV56		10	13		HPV56		9	408 173
HPV56		11	4		HPV56		11	246
HPV56		8	467		HPV56		8	420
HPV56		11	467 442		HPV56		10	87
HPV56			52		HPV56		9	214
HPV56 HPV56		11	494		HPV56		11	214
HPV56		9	494		HPV56		10	367
HPV56		10	494		HPV56		10	324
HPV56		9	406		HPV56		11	324
HPV56		11	406		HPV56		9	281
HPV56		11	189		HPV56		10	281
HPV56		8	288		HPV56		8	7
HPV56		8	339		HPV56		10	7
HPV56		9	339		HPV56	L1	9	423
HPV56		10	384		HPV56		10	423
HPV56		10	213		HPV56		9	268
HPV56		9	395		HPV56		8	396
HPV56		11	395		HPV56		10	396
HPV56		9	103		HPV56		8	283
HPV56		10	159		HPV56		9	325
HPV56		9	76		HPV56	L1	10	325
HPV56		10	76		HPV56	L1	8	62
HPV56		8	110		HPV56	L1	9	62

Table X HLA A24 Supermotif Peptides

			*****	oupermous	· r optide	•		
HPV56	L2	9	240		HPV56	L2	8	235
HPV56		11	286		HPV56	L2	9	255
HPV56		8	438		HPV56		11	255
HPV56		11	303		HPV56	L2	8	161
HPV56		10	246		HPV56		8	50
HPV56		11	246		HPV56		10	50
HPV56		9	367		HPV56		9 .	181
HPV56		10	14		HPV56		8	337
HPV56		10	201		HPV56		11	337
HPV56		8	275		HPV56		8	106
HPV56		11	322		HPV56		8	248
HPV56		11	406		HPV56		9	248
HPV56		8	425		HPV56		8	347
HPV56		9	83		HPV56		8	121
HPV56		11	30		HPV56		10	121
HPV56		9	429		HPV56		9	353
HPV56		11	429		HPV56		11	179
HPV56		8	331		HPV56		10	278
HPV56			398		HPV56		9	385
		10			HPV56		10	388
HPV56		8	175		HPV56		8	395
HPV56		10	444		HPV56		8	417
HPV56		8	241		HPV56		-	400
HPV56		9	122		HPV56		8 10	400
HPV56		10	287					325
HPV56		9	51		HPV56		8 10	325
HPV56		9	401		HPV56			
HPV56		10	217		HPV56		9	374
HPV56		11	217		HPV56		8	214
HPV56		8	188		HPV56			209
HPV56		10	188		HPV56		10	254 160
HPV56		11	118		HPV56		9	
HPV56		9	346		HPV56		10	392
HPV56		11	25		HPV56		11	392
HPV56		8	258		HPV56		9	73
HPV56		10	206		HPV56		9	336
HPV56		10	62		HPV56		9	98
HPV56		10	310		HPV56		9	410
HPV56		11	310		HPV56		8	185
HPV56		8	269		HPV56		11	185
HPV56		9	269		HPV56		9	423
HPV56		11	269		HPV56		10	423
HPV56		8	156		HPV56		8	312
HPV56		9	372		HPV56		9	312
HPV56		11	372		HPV56		10	312
HPV56		11	151		HPV56		8	16
HPV56		9	318		HPV56		9	172
HPV56		10	180		HPV56		10	172
HPV56		8	44		HPV56		11	172
HPV56		10	44		HPV56		8	306
HPV56		11	44		HPV56		9	233
HPV56		9	279		HPV56		10	233
HPV56		11	81		HPV56		8	46
HPV56		10	407		HPV56		9	46
HPV56		11	103		HPV56		11	295
HPV56		11	34		HPV56		8	220
HPV56		8	43		HPV56		8	298
HPV56		9	43		HPV56		8	316
HPV56		11	43		HPV56		11	316
HPV56		8	38		HPV56		8	436
HPV56	L2	10	38		HPV56	L2	9	436

Table X HLA A24 Supermotif Peptides

HPV56	L2	10	436
HPV56	L2	8	343
HPV56	L2	9	343
HPV56	L2	9	49
HPV56	L2	11	49
HPV56	L2	8	154
HPV56	L2	9	154
HPV56	L2	10	154
HPV56	L2	10	212
HPV56	L2	10	183
HPV56	L2	8	134
HPV56	L2	10	134
HPV56	L2	10	148
HPV56	L2	11	414
HPV56	L2	9	365
HPV56	L2	11	365
HPV56	L2	9	95
HPV56	L2	9	111
HPV56	L2	10	191
HPV56	L2	10	304
HPV56	L2	9	105
HPV56	L2	9	247
HPV56	L2	10	247
HPV56	L2	8	386
HPV56	L2	8	136
HPV56	L2	9	288
HPV56	L2	9	135
HPV56	L2	9	149
HPV56	L2	9	39
HPV56	L2	10	415
HPV56	L2	8	52
HPV56	L2	8 .	112
HPV56	L2	11	112
HPV56	L2	9	408
HPV56	L2	11	408
HPV56	L2	9	389
HPV56	L2	10	104
HPV56	L2	8	84
HPV56	L2	10	31
HPV56	L2	8	430
HPV56	L2	10	430
HPV56	L2	11 .	430
HPV56	L2	11	443
HPV56	L2	9	431
HPV56	L2	10	431
HPV56	L2	11	71
HPV56	L2	8	319

2	3	4	E2	8	35
E1	10	377	E2	9	35
E1	11	392	E2	10	35
				8	488
L2	9	238	E1		
L2	10	238	L1	9	153
L2	8	275	E1	10	14
E1	9	251	E6	8	131
E1	11	251	E6	11	131
E5	9	23	E6	8	31
E5	10	23	E1	10	640
E5	11	23	E4	9	64
L2	8	286	E1	8	529
	11	286	E1	9	529
L2				9	140
E2	8	72	E6		
E2	11	72	E6	10	140
L2	10	112	E5	8	75
E1	11	22	E5	11	75
E4	8	14	E6	8	104
			E1	9	385
E4	9	14			
E4	11	14	E1	10	385
E1	11	520	E1	8	49
E1	10	207	E5	9	39
Li	8	81	E5	11	39
			E1	8	369
L2	8	421			
L2	9	421	E1	10	369
E1	10	554	L1	9	219
E1	11	554	L2	10	278
E6	8	37	E6	10	96
E6	10	37	E6	11	96
			E7	8	75
E5	8	79	E7	9	75
E1	9	319			
L1	11	22	E7	10	75
E1	11	296	L2	9	404
E4	9	61	E1	8	570
L2	10	14	E1	10	570
	8	525	E1	11	570
E1					347
E1	9	525	L2	11	
E6	11	10	L2	10	396
E1	10	77	E2	9	313
E1	11	77	E1	11	81
E1	10	101	E2	9	25
			Li	8	366
L1	9	43	Li		366
E1	10	601		11	
E1	9	271	E7	10	14
L1	11	384	L1	11	208
E6	8	47	E1	10	203
E6	9	47	E1	11	46
L1	8	25	L1	10	195
					141
L1	9	25	E2	11	
L1	10	25	L2 '	10	344
E1	10	612	L1	9	198
E1	9	215	E1	9	481
E1	10	215	E1	11	481
			E1	10	73
E5	8	27			
E5	10	27	L1	8	331
E5	11	27	L1	10	331
E6	8	67	E1	8	534
E6	11	67	L1	11	411
E6	9	137	L2	11	30
	9	296	E6	8	99
E2	7	490	10		23

			TIEA-A2+ Supermon	i i cpiide	3	
L2	11	258		L1	10	369
E2	10	136		E6	8	126
E1	9	236		E1	8	454
L1	9	146		E1	10	454
L1	10	229		E1	11	454
E2	9	130		L2	8	428
E6	٠. و	69		L2	9	428
E1	9	453		E5	8	40
El	11	453		E5	10	40
L2	9	192		E5	9	68
E1	و	105		E1	10	393
E1	11	105		E2	11	346
L2	10	120		L2	8	398
E6	10	42		L2	10	398
L1	11	453		E1	9	446
E1	8	197		E1	10	446
E1	10	604		L1	8	245
E1	11	604		E1	8	457
E1	9	131		E1	9	457
E2	10	17		L2	8	239
E2	9	74		L2	9	239
E2	10	74		L2	11	239
E1	10	417		E1	9	587
E1	11	417		E2	8	171
E1	11	360		E5	9	28
E2	11	100		E5	10	28
E7	10	73		B5	8	24
E7	11	73		E5	9	24
E6	10-	92		E5	10	24
E6	11	92		E5	11	24
L2	8	135		L1	10	441
E4	9	75		L1	9	33
E2	9	185		L1	10	33
E7	10	39		L2	8	434
E1	9	39		L2	9	434
E6	8	113		L1	10	200
E6	9	113		E1	8	241
L1	10	262		E2	8	361
L1	11	262		L2		433
L1	8	103		L2	9	433
L1	9	381		L2	10	433
Li	8	443		L1	8	318
E7	10	78		L1	9	318
E2	10	205		E1	10	243
E1	9	339		E1	11	194
E1	8	379		E1	9	326
Ll	8	364		E2	9	156
L1	10	364		B1	9	350
L1	9	357		Li	10	101
L1	11	357		E7	8	22
E2	9	86		E1	8	217
E1	9	613		E1	11	217
E1	11	613		L1	10	400
L1	9	243		L2	8	292
L1	10	243		L1	9	144
E5	10	67		L1	11	144
L1	8	244		E2	8	55
L1	9	244		E1	10	273
	8	220		E1	8	191
E1 L1	8	369		E1	9	191
n T	٠	300			-	

			rila-A24 Superinoni	repudes		
L2	10	215		E7	11	5
L2	11	215		E2	11	32
L2	8	25		E1	8	75
L2	8	64		L2	9	318
L2	11	64		E1	10	412
E1	8	436		L1	9	370
E1	11	436		L1	10	32
E7		85		L1	11	32
	9	407		E5	8	21
L1		413		E5	11	21
L2	10			E5	10	31
E1	8	467 467		E5	11	31
E1	10			L2	9	113
L2	8	75			9	279
L1	9	222		L2 E6	9	97
L1	10	222		E6	10	97
E5	8	11		E1	10	195
E5	11	11				178
E4	8	90		L2	8	
E4	10	90		E5	9	32
E1	8	316		E5	10	32
L2	9	51		L2	10	44
L1	9	111		L2	11	44
L1	10	111		E5	8	17
Ē2	10	242		E6	8	120
L1	8	113		E6	10	120
L1	11	113		L2	8	405
E1	8	16		L2	8	429
E1	8	44		E1	11	56
E6	11	58		E1	9	571
L1	11	64		E1	10	571
L1	8	468		L1	8	376
L2	8	312		L1	10	376
L2	9	312		L1	11	376
L2	10	312		E1	11	341
E2	10	53		L2	9	185
L2	9	177		L2	11	185
E6	8	119		L2	11	131
E6	9	119		L1	10	187
E6	11	119		L2	8	247
E1	9	264		E1	11	476
E1	11	264		L2	9	121
E4	8	59		L2	11	121
E4	11	59		E1	10	443
E2	10	78		E2	11	287
E2	10	310		E1	10	23
E4	10	10		L2	10	104
E6	9	25		L2	11	104
L1	8	387		E5	8	34
L1	11	387		E5	10	34
E4	10	26		E5	9	41
L2	8	306		E5	11	41
E1	9	581		L2	10	348
L1	9	361		E1	10	531
Ll	10	361		E1	11	531
L1	11	361		E6	9	43
		29		L1	10	286
E2	8	502		E2	8	157
E1	8	502		E2	11	157
E1	9			L1	9	79
E1	10	502		L1	10 .	79
E7	9	5		-1	-0 .	, ,

Table XA HPV6A HLA-A24 Supermotif Peptides

			IILA-A24 Superino	штери	ues	
E2	10	120		E1	8	557
E1	8	211		E5	9	16
El	10	211		E6	11	101
E1	11	211		E1	10	491
L1	8	462		E2	8	314
L1	9	449		L1	11	186
L1	11	449		L2	9	246
E1	8	433		E5	8	33
E1	11	433		E5	9	33
E6	8	73		E5	11	33
E6	10	73		L1	8	41
E2	10	351		L1	9	41
E1	9	312		L1	11	41
E1	10	312		E1	10	521
E1	11	312		E1	11	521
E2	9	359		E1	9	540
E2	10	359		E1	9	208
E1	8	254		E1	11	208
E1	9	254		E7	11	83
E6	11	128		E4	8	8
E1	8	357		E1	11	198
E1	10	357		E2	8	82
L2	11	22		E2	11	82
E1	8	420		E7	8	82
E2	9	84		E5	9	59
E2	11	84		E5	10	59
E6	10	18		E5	11	59
E4	8	42		E5	11	55
L1	10	56		E5	8	51
L2	11	34		E5	9	51
E2	11	147		E5	10	51
E6	11	116		E5	11	51
E6	8	52		E1	9	298
E6	10	52		E1	10	298
E2	11	63		E1	11	298
L1	10	61		E5	8	69
L1	8	19		E5 *	8	60
L1	10	71		E5	9	60
L2	8	46		E5	10	60
L2	9	46		E5	11	60
E2	8	177		E5	9	72
E1	11	85		E5	10	72
E1	10	578		E5	11	72
L1	8	226		E1	9	563
E4	8	18		E5	10	56
E1	9	401		E2	8	42
E2	9	311		E2	10	42
E2	11	311		E5	8	52
L1	9	87		E5	9	52
L1	11	87		E5	10	52
L1	10	242		E1	8	132
L1	11	242		E1	9	358
L2	9	397		L2	10	23
L2	11	397		E6	9	121
L1	8	302		E1	9	555
E1	11	66		E1	10	555
E6	8	54		E5	8	49
L2	8	107		E5	9	49
E1	8	255		E5	10	49
E1	11	255	•	E5	11	49

				p		
E5	11	70		L2	8	38
E1	9	268		L2	10	38
E1	10	268		E1	11	96
E6	9	38		E2	8	127
E6	11	38		L2 ·	9	385
E5	8	61		L1	11	142
E5	9	61		E2	9	348
E5	10	61		E1	9	609
L2	9	338		E1	8	439
L2	11	338		E1	9	439
E5	8	73		E1	10	594
E5	9	73		E1	8	456
E5	10	73		E1	9.	456
E1	8	514		E1	10	456
E1	9	514		L2	8	415
E5	8	47		L2	9	325
E5	9	47		L2	11	325
E5	10	47		L1	11	217
E5	11	47		L2	10	189
L1	8	295		E1	11	442
L1	9	295		E6	11	110
E1	8	564		L1 L1	9	183 458
E7	10	67		L2	10	73
L1	8	95 57		L1	11	109
E5	9	57		E7	8	47
E5 E1	11 8	261		E1	8	562
E2	9	18		E1	10	562
E2	9	43		E5	9	64
E4	9	11		E5	10	64
E4	11	11		E1	8	258
E5	8	53		E1	10	258
E5	9	53		E1	11	258
E4	8	5		L2	9	389
E4	9	5		L2	10	389
E4	11	5		L2	11	389
E1	8	320		L2	10	337
E1	8	306		E1	9	513
E2	8	151		E1	10	513
E2	9	151		E1	11	545
E1	10	47		L2	9	408
L1	9	196		E2 E2	10	354 354
L1	11	196		L2	8	183
L1	8	154		L2	11	183
E1 E1	9 10	274 361		L1	11	426
E2	10	101		L2	9	359
L1	10	1		Li	8	90
Ē2	9	170		L2	9	96
E1	10	568		E2	9	258
E1	8	451		E2	10	258
E1	11	451		L2	8	171
L1	11	31		L2	9	171
E1	8	300		L2	10	171
E1	9	300		L2	9	426
L1	10	445		L2	10	426
E1	8	539		L2	11	426
El	10	539		E7	10	20
L1	9	438		E5	10	19
E6	9	21		L1	8	266

			IILA-A24 Superillou	i repude	•	
Ll	10	175		E1	11	266
L1	11	16		L1	8	456
L2	11	418		L1	9	456
L2	9	363		L1	10	456
L2	10	363		E6	10	76
L2	9	382		L1	9	66
E2	9	91		E6	9	125
E2	11	91		L2	8 ·	298
E4	11	71		L2	8	316
E6	9	60 60		L2 E2	11 9	316 7
E6	11	237		E2 E2	10	7
L1 L1	9	237		L1	11	241
L1	10	237		E1	8	125
L1	8	435		L2	10	245
E1	9	90		L1	9	40
E5	8	66		L1	10	40
E5	11	66		E2	9	303
L1	9	368		E2	10	303
L1	11	368		E1	8	616
E1	10	355		E7	11	66
E2	9	163		L1	9	94
E2	8	345		E1	11	324
E1	8	445		E1	9	293
E1	10	445		L1 L1	9	279 279
E1	11	445 289		L2	9	221
E2 E5	9 10	7		L1	8	140
E5	11	7		E1	11	583
L2	8	43		E2	11	301
L2	11	43		L2	11	295
E6	11	28		L1	8	414
E6	8	15		E1	9	219
E6	10	15		E1	8	493
L1	11	151		L1	11	440
L1	8	301		E1	9 .	547
L1	9	301		E1	11	547
E6	10	50		L2	8	153
E7	9	81		L2 L2	10 11	267 267
L2	8	16 4		ES	8	30
E4 E4	9	4		E5	11	30
E4	10	4		E2	8	207
L1	11	232		L2	8	392
L1	8	250		E1		247
E2	9	48		E1	10	247
E2	8	76		L1	9	375
E1	9	305		L1		375
L1	9	210		L2	11	103
L1	11	210		L1	11	285
E2	8	103		E1	8	60
E1	10	128		L1	10	86
E1	8	344		L2 L2		49 49
L2 L2	9 10	231 231		L2		106
L2 L2	8	231		L2		106
E1	8	363		E1		490
E2	11	183		E2		81
E1	10	425		L1		294
E1	9	266		L1		294

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E1	8	260	L1	8	408
E1	9	260	E1	8	556
E6	11	23	E1	9	556
L2 '	10	304	E5	8	15
E2	8	150	E5	10	15
E2	9 .	150	E7	9	7
E2	10	150	E5	8	50
E2	9	23	E5	9	50
E2	11	23	E5	10	50
E2	9	180	E5	11	50
E5	8	14	E1	10	297
E5	9	14	E1 E5	11 10	297
E5	11	374	E5	11	71 71
L2 L2	10 9	241	E2	9	93
E6	10	7	E2	10	93
E2	8	201	E6	8	26
E1	10	289	L1	9	377
El	11	289	Li	10	377
E1	10	331	E1	11	408
E1	11	331	E2	11	128
L1	8	348	L1	10	388
L1	11	348	E6	8	39
L1	8	189	E6	10	39
L1	8	392	E1	11	232
L1	10	392	L1	8	223
E2	10	40	L1	9	223
E5	8	45	L1	11	223
E5	9	45	E6	8	142
E5	10	45	E1	9	585
E5	11	45	E1	11	585
L2	9	260	E6	10	11
E1	11	185	E5	8	62
E2	11	198	E5	9	62
L2	10	145	E5	11	62
L2	11	145	L1	9	332
L1	8	343	L2	10	151
E2	8	144	L1	11	345
E6	10	4.5	E5	10	12
E6	11	45 485	E5 L2	11	12 150
E1 E1	8 10	485	E4	8	76
E1	11	485	E6	11	87
L2	9	345	L2	8	386
E6	9	19	E4	9	91
E6	11	19	E1	9	290
E1	8	588	E1	10	290
L2	11	406	E6	10	88
E1	8	586	E6	11	88
E1	10	586	E4	9	73
L2	9	39	E4	11	73
E1	10	97	E1	9	332
E6	9	12	E1	10	332
E6	11	12	L2	11	387
E2	8	355	E1	9	78
E2	9	355	E1	10	78
E2	11	355	E4	8	92
L2	10	184	L1	11	328
E1	8	294	L1	9	57
E7	8	86	L1	11	57

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E6	10	132			E2	9	137
E2	11	336			E1	9	426
L1	10	412			E5	8	36
L1	10	349			E1	8	340
L1	11	349			E1	8	530
L2	8	52			E1	11	530
L2	8	427			E1	11	464
L2	9	427			E5	8	58
L2	10	427			E5	10	58
E1	11	346			E5	11	58
E1	8	333			E1	8	267
E1	9	333			E1	10	267
E5	9	20			E1	11	267
L2	10	31			E2	8	92
E1	11	499			E2	10	92
L1	9	393			E2	11	92
E6	9	53			E6	8	141
E4	9	7			E6	9	141
E1	8	275			E4	10	72
E2	9	41			E1	8	237
E2	11	41			L2	11	440
L1	8	73			L2	10	441
L1	10	73			E1	9	486
E5	8	48			E1	10	486
E5	9	48			L2	8	319
E5	10	48			L1	10	385
E5	11	48			E4	8	12
E5	8	46			E4	10	12
E5	9	46			E4	11	12
E5	10	46			E1	10	86
E5	11	46			L2	8	435
L1	8	382			E1	9	579
L2	10	419			E1	11	579
L2	11	419			E5	8	54
E7	8	6			L1	9	230
E7	10	6			E1	9	532
L2	8	364			E1	10	532
L2	9	364			L1	8	358
L1	8	27			L1	10	358
L2	9	146			E1	11	536
L2	10	146			E2	9	159
E1	10	584			E2	10	159
E1	11	238			L1	9	350
E5	8	25			L1	10	350
E5	9	25			E5	9	43
E5	10	25			E5	10	43
L1	10	329			E5	11	43
E2	8	349			E1	8	402
L1	9	72			E4	8	6
L1	11	72			E4	10	6
L1	8	58			E2	11	168
L1	10	58			L1	9	287
E7	9	68			E2	8	138
E5	9	35			E1	8	91
E4	8	65			L1	8	26
L2	11	253			Ll	9	26
E1	9	602			E2	8	131
E5	8	42			E2	10	158
E5	10	42			E2	11	158
E5	11	42					

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2	3	4		E1	9	234
El	10	377		E1 E6	11 8	234 67
E5B	9	21		E6	11	67
E5B	10	21		E6	9	
E5B	11	21			9	137
E1	11	392		E2 E2	. 8	296 35
L2	9	238		E2 -	9	35
L2	10	238		E2	10	35
L2	8	275		E1	8	488
E1	9	251		L1	9	153
E1	11	251		E1	10	14
E5A E5A	9 10	23 23		E6	8	131
	11	23		E6	11	131
E5A L2	8	286		E6	8	31
L2	11	286		E1	10	640
E2	8	72		E4	9	74
E2	11	72		E1	8	529
L2	10	112		E1	9	529
E1	11	22		E6	9	140
E4	8	24		E6	10	140
E4	9	24		E5A	8	75
E4	11	24		E5A	10	75
E1	11	520		E5A	11	75
E1	10	207		E6	8	104
L1	8	81		E1	9	385
L2	8	421		E1	10	385
L2	9	421		E1	8	49
E1	10	554		E1	- 8	369
E1	11	554		E1	10	369
E6	8	37		L1	9	219
E6	10	37		L2	10	278
ESA	8	79		E6	10	96
E1	9	319		E6	11	96
Ll	11	22		E7	8	75
E1	11	296		E7 E7	10	75 75
E4	9	71		L2	9	403
L2 E1	10 8	14 525		E1	8	570
E1	9	525		El	10	570
E6	11	10		E1	11	570
E1	10	77		L2	11	347
E1	11	77		L2	10	395
E1	10	101		E2	9	25
Li	9	43		E2	9	313
E1	10	601		E1	11	81
E1	9	271		L1	8	366
E1	10	271		L1	11	366
L1	11	384		E7	10	14
E6	8	47		L1	11	208
E6	9	47		E1	10	203
L1	8	25		E1	11	46
L1	9	25		L1	10	195
L1	10	25		L2	10	344
El	10	612		Ll	9	198
E1	9	215		E1	9	481
El	10	215		E1	11	481
E5A	8	27		E1	10	73
E5A	10	27		L1 L1	8 10	331 331
E5A	11	27		PT	10	221

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E5B	8	11		E1	9	613
E5B	9	11		E1	11	613
E2	9	143		L1	9	243
E1	8	534		Ll	10	243
L1	11	411		E5A	10	67
L2	11	30		L1	8	244
E6	8	99		L1	9	244
L2	11	258		E1	8	220
E2	9	348		L1	8	369
E2	10	136		L1	10	369
El	9	236		E6	8	126
Li	9	146		E5A	9	16
L1	10	229		E1	8	454
E2	9	130		E1	10	454
E6	9	69		E1	11	454
E1	9	453		L2	8	428
E1	11	453		L2	9	428
E5A	9	39		E5B	8	22
E5A	11	39		E5B	9	22
E1	9	105		E5B	10	22
E1	11	105		E5B	11	22
L2	10	120		E5A	8	40
E6	10	42		E5A	10	40
L1	11	453		E2	11	346
E1	8	197		E5A	9	68
E1	10	604		E1	10	393
E1	11	604		L2	8	397
E1	9	131		L2	10	397
E2	10	17		E1	9	446
E2	9	74		E1	10	446
E2	10	74		L1	8	245
E1	10	417		L2	8	239
E1	11	417		L2	9	239
E1	11	360		L2	11	239
E2	11	100		E1	8	457
E7	10	73		E1	9	457
E7	11	73		E1	9	587
E6	10	92		E2	8	171
E6	11	92		E5A	9	28
L2	8	135		E5A	10	28
E4	9	85		E5A	8	24
E2	9	185		E5A	9	24
E7	10	39		E5A	10	24
E1	9	39		E5A	11	24
E6	8	113		L1	10	441
E6	9	113		L1	9	33
L1	10	262		L1	10	33
L1	11	262		L2	8	434
L1	8	103		L2	9	434
L1	9	381		L1	10	200
L1	8	443		B1	8	241
E7	10	78		E2	8	361
E2	10	205		L2	8	433
E1	9	339		L2	9	433
E1	8	379		L2	10	433
L1	8	364		L1	8	318
L1	10	364		L1	9	318
L1	9	357		E1	10	243
L1	11	357		E1	11	194
E2	9	86		E1	9	326

			HLA-A24 Supermon	i Peptide	s	
E2	9	156		E6	10	50
E1	9	350		E4	10	20
Li	10	101		E6	9	25
E7	8	22		L1	8	387
B1	8	217		L1	11	387
E1	11	217		E4	10	36
L1	10	400		L2	8	306
L2	8	292		E1	9	581
E5B	8	15		L1	9	361
E5B	9	15		L1	10	361
L1	9	144		L1	11	361
L1	11	144		E2	8	29
E5B	8	25		E1	8	502
E5B	10	25		E1	9	502
E2	8	55		E1	10	502
E1	8	273		E7	9	5
E1	10	273		E7	11	5
E1	8	191		E2	11	183
E1	9	191		E2	11	32
L2	10	215		E1	8	75
L2	11	215		L2	9	318
L2	8	25		E1	10	412
L2	8	64		E5B	8	61
L2	11	64		E5B	9	61
E1	8	436		E5B	10	61
E1	11	436		E5B	11	61
L1	9	407		L1	9	370
E7	9	85		L1 L1	10 11	32 32
L2	11 8	4·12 467		E5A	8	21
E1 E1	10	467		E5A	11	21
Ll	9	222		E5A	10	31
Ll	10	222		E5A	11	31
ESA	8	11		L2	9	113
E5A	11	11		L2	9	279
E4	8	100		E6	9	97
E4	10	100		E6	10	97
E1	8	316		E1	10	195
L2	9	51		E5A	9	32
L1	9	111		E5A	10	32
L1	10	111		L2	10	44
E2	10	242		L2	11	44
L1	8	113		E5A	8	17
L1	11	113		E6	8	120
E1	8	16		E6	10 8	120
E1	8	44		L2 L2	8	404 429
E6 L1	11 11	58 64		E1	11	56
L1	8	468		E1	9	571
L2	8	312		E1	10	571
L2	9	312		Li	8	376
L2	10	312		Li	10	376
E2	10	53		L1	11	376
E6	8	119		E1	11	341
E6	9	119		L2	11	131
E6	11	119		L1	10	187
E1	9	264		L2	8	247
E1	11	264		E5B	8	23
E2	10	78		E5B	9	23
E2	10	310		E5B	10	23

			Table 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	rreplice		
E1	11	476		E1	11	85
L2	9	121		E1	10	578
L2	11	121		Ll	8	226
E1	10	443		E4	8	28
E2	11	287		E1	9	401
E1	10	23		E4	8	8
L2	10	104		E4	9	8
L2	11	104		E4	11	8
E5A	8	34		E2	9	311
E5A	10	34		E2	11	311
E5A	9	41		L1	9	87
E5A	11	41		L1	11	87
L2	10	348		L1	10	242
E1	10	531		L1	11	242
E1	11	531		L2	9	396
E6	9	43		L2	11	396
Li	10	286		L1	8	302
E2	8	157		E1	11	66
E2	11	157		E6	8	54
Ll	9	79		L2	8	107
LI	10	79		E1	8	255
E2	10	120		E1	11	255
EÍ	8	211		E1	8	557
E1	10	211		E6	11	101
E1	11	211		E1	10	491
L1	8	462		E2	8	314
Ll	9	449		Li	11	186
Ll	11	449		L2	9	246
E1	8	433		E5A	8	33
E1	11	433		E5A	9	33
E6	8	73		E5A	11	33
E6	10	73		Ll	8	41
E2	10	351		L1	9	41
E1	9	312		Li	11	41
E1	10	312		E5B	9	18
E1	11	312		E5B	10	18
E2	9	359		E1	10	521
E2	10	359		E1	11	521
E1	8	254		E1	9	540
E1	9	254		El	9	208
E6	11	128		E1	11	208
E1	8	357		E7	11	83
E1	10	357		E4	8	18
L2	11	22		E1	11	198
E1	8	420		E7	8	82
E6	10	18		E5A	9	59
E4	8	52		E5A	10	59
L1	10	56		E5A	11	59
L2	11	34		E5A	11	55
E2	11	147		E5A	8	51
E6	11	116		E5A	9	51
E6	8	52		E5A	10	51
E6	10	52		E5A	11	51
E2	11	63		E1	9	298
L1	10	61		E1	10	298
L1	8	19		E1	11	298
L1	10	71		E2	8	82
L2				E2	11	82
	8	46				
L2 E2	8 9 8	46 17 7		E5A E5A	8	69 60

Table XB HPV6B HLA-A24 Supermotif Peptides

			IILA-A24	Supermon	t i cpuo			
E5A	9	60			E4	8	15	
E5A	10	60			E4	9	15	
E5A	11	60			E4	11	15	
E5A	9	72			E1	8	320	
E5A	10	72			E5B	9	26	
E5A	11	72			E1	8	306	
E1	9	563			E2	8	151	
E5A	10	56			E2	9	151	
E2	8	42			E1	10	47	
E2	10	42			L1	9	196	
E5A	8	52			Ll	11	196	
E5A	9	52			L1	8	154 274	
E5A	10	52			E1	9		
E5B	11	3			E1	10 10	361 101	
E1	8	132			E2 L1	10	1	
E1	9	358			E2	9	170	
L2	10	23			E1	10	568	
E6	9	121			E1	8	451	
E1	8	458			E1	11	451	
E1	9	555			Ll	11	31	
E1	10	555 49			E4	11	5	
E5A	8 9	49			El	8	300	
E5A	10	49			E1	9	300	
E5A E5A	11	49			Li	10	445	
E5A	11	70			E1	8	539	
E1	9	268			E1	10	539	
E1	10	268			L1	9	438	
E6	9	38			E6	9	21	
E6	11	38			L2	8	38	
E5A	8	61			L2	10	38	
E5A	9	61			E1	11	96	
E5A	10	61			E2	8	127	
L2	9	338			E1	8	607	
L2	11	338			E1	11	607	
E5A	8	73			L2	8	385	
E5A	9	73			L1	11	142	
ESA	10	7.3			. E1	9	609 439	
E1	8	514			E1	9	439	
E1	9	514			E1 E1	10	594	
E5A	8	47			El	10	226	
E5A	9	47			E1	8	456	
E5A	10	47			E1	9	456	
E5A	11	47			E1	10	456	
L1	8	295 295			L2	9	414	
L1	9	564			L2	9	325	
E1 E7	8 10	67			L2	11	325	
L1	8	95			L1	11	217	
ESB	8	16			L2	10	189	
E5B	11	16			E1	11	442	
E5A	9	57			E4	8	69	
E5A	11	57			E4	11	69	
E1	8	261			E6	11	110	
E2	9	18			L1	9	183	
E2	9	43			L1	8	458	
E4	9	21			L1	11	109	
E4	11	21			E7	8	47	
ESA	8	53			E1	8	562	
E5A	9	53			E1	10	562	

E5A	9	64		ESB	9	6
E5A	10	64		ESB	11	6
L2	10	73		E2	9	289
				E5A	10	7
L2	8	389		E5A	11	7
L2	9	389		L2	8	43
L2	10	389				43
E1	8	258		L2	11	
E1	10	258		E6	11	28
E1	11:	258		E6	8	15
L2	10	337		E6	10	15
E1	9	513		L1	11	151
E1	10	513		L1	8	301
E1	11	545		L1	9	301
L2	9	407		E7	9	81
E2	9	354		L2	8	16
E2	10	354		E4	8	14
		426		E4	9	14
L1	11			E4	10	14
L2	9 .	359		L1	11	232
L1	8	90		Li	8	250
L2	8	183				
L2	11	183		E2	9	48
L2	9	96		E2	8	76
E2	9	258		E1	9	305
E2	10	258		L1	9	210
L2	8	171		L1	11	210
L2	9	171		E2	8	103
L2	10	171		E1	10	128
L2	9	426		L2	8	75
L2	10	426		E1	8	344
L2	11	426		L2	9	231
	10	20		L2	10	231
E7		19		L2	8	233
E5A	10			E1	8	363
L1	8	266		E1	10	425
L1	10	175		E1	9	266
L1	11	16		E1	11	266
L2	9	363				456
L2	10	363		L1	8	
L2	9	382		L1	9	456
L2	11	382		L1	10	456
E2	9	91		E6	10	76
E2	11	91		L1	9	66
E4	11	81		E6	9	125
E6	9	60		L2	8	298
E6	11	60		L2	8	316
L1	8	237		L2	11	316
L1	9	237		E2	9	7
Ll	10	237		E2	10	7
L1	8	435		L1	11	241
				E1	8	125
E1	9	90		L2	10	245
ESA	8	66		L1	9	40
E5A	11	66		L1	10	40
L1	9	368			9	303
L1	11	368		E2		303
E1	10	355		E2	10	
E2	9	163		E1	8	616
E2	8	345		E7	11	66
El	8	445		Ll	9	94
E1	10	445		E2	9	84
E1	11	445		E2	11	84
E5B	8	6		E1	11	324
220	3	•				

E1	9	293		L1	8	189
Ll	8	279		L1	8	392
L1	9	279		L1	10	392
L2	9	221		E2	10	40
L1	8	140		E5A	8	45 45
E1	11	583		E5A E5A	9 10	45
E2	11	301			11	45
L2	11	295		E5A L2	9	260
L1	8	414		E1	11	185
E1	9	219		E2	11	198
E1	8	493 440		L2	10	145
L1	11 9	547		L2	11	145
E1	11	547		Li	8	343
E1 L2	8	153		E6	10	45
L2	10	267		E6	11	45
L2	11	267		E1	8	485
E5A	8 _	30		E1	10	485
E5A	11	30		E1	11	485
E2	8	207		L2	9	345
E1	9	247		E5A	8	15
E1	10	247		E5A	10	15
L1	9	375		E6	9	19
L1	11	375		E6	11	19
L2	11	103		E1	8	588
L1	11	285		L2	11	405 586
E1	8	60		E1	8 10	586
L1	10	86		E1 L2	9	39
L2	9	49		E1	10	97
L2	11	49		E6	9	12
L2	8	106		E6	11	12
L2	9	106		E2	8	355
E1	11	490 81		E2	9	355
E2	9 8	391		E2	11	355
L2 L1	9	294		E1	8	294
L1	10	294		L1	8	408
E1	8	260		E1	8	556
E1	9	260		E1	9	556
E6	11	23		E7	9	7
L2	10	304		E5A	8	50
E2	8	150		E5A	9	50
E2	9	150		E5A	10	50
E2	10	150		E5A	11	50
E2	9	23		E1	10	297 297
E2	11	23		E1	11	71
E5A	8	14		E5A E5A	10 11	71
E5A	9	14		E5A E7	8 11	86
E5A	11	14		E2	9	93
E2	9	180		E2	10	93
L2	10	374		E6	8	26
L2	9	241		L1	9	377
E6	10	7		L1	10	377
E2	8	201 289		E1	11	408
E1	10	289		E2	11	128
E1	11	331		E5B	10	59
E1 E1	10 11	331		E5B	11	59
L1	8	348		L1	10	388
L1	11	348		E6.	8	39
11.1		5				

E6	10	39	E1	11	499
L1	8	223	L1	9	393
L1	9	223	E6	9	53
L1	11	223	E4	9	17
E1	11	232	E1	8	275
E6	8	142 .	E2	9	41
E5B	8	63	E2	11	41
E5B	9	63	L1	8	73
E1	9 '	585	L1	10	73
E1	11	585	ESA	8	48
E6	10	11	E5A	9	48
ESA	8	62	E5A	10	48
E5A	9	62	E5A	11	48
E5A	11	62	E5A	8	46
L1	9	332	E5A	9	46
L2	10	151	E5A	10	46
L1	11	345	E5A	11	46
E5A	10	12	E4	9	10
ESA	11	12	E4	10	10
L2	11	150	L1	8	382
E4	8	86	E7	8	6
E6.	11	87	E7	10	6
L2	11	386	E5B	11	58
E4	9	101	L2	8	364
E2	11	212	L2	9	364
E1	9	290	L2	11	418
E1	10	290	L1	8	27 185
E6	10	88	L2	9	185
E6	11	88	L2	11	
E4	9	83	L2 L2	9 10	146 146
E4	11	83		10	584
E1	9	332	E1 E1	11	238
E1	10	332	E5A	8	25
L2	10	387	ESA	9	25
L2	11	387	ESA	10	25
E1	9	78	L1	10	329
E1	10	78	E2	8	349
L2	10	184	L1	9	72
E4	8	102	L1	11	72
L1	11	328	L1	8	58
L1	9 11	57 57	L1	10	58
L1	10	132	E7	9	68
E6	8	132	E5A	9	35
E5B E5B	11	12	E4	8	75
E2	8	144	L2	11	253
E2	11	336	E1	9	602
L1	10	412	ESA	8	42
L1	10	349	E5A	10	42
L1	11	349	E5A	11	42
E2	10	213	E2	9	137
L2	8	52	E1	9	426
L2	8	427	E5A	8	36
L2	9	427	E1	8	340
L2	10	427	E1	8	530
E1	11	346	E1	11	530
E1	8	333	ESB	10	13
E1	9	333	E5B	11	13
E5A	9	20	E1	11	464
L2	10	31	E5B	10	17

E5B	11	17
		58
E5A	8	
E5A	10	58
E5A	11	58
E1	8	267
E1	10	267
E1	11	267
	8	92
E2		
E2	10	92
E2	11	92
E6	8	141
E6	9	141
E4	-10	82
E1	8	237
L2	11	440
		441
L2	10	
E1	9	486
E1	10	486
L2	8	319
L1	10	385
E4	8	22
E4	10	22
		22
E4	11	
E1	10	86
L2	8	435
E1	9	579
E1	11	579
E5A	8	54
L1	9	230
	é	532
E1		532
E1	10	
L1	8	358
L1	10	358
E1	11	536
E2	9	159
E2	10	159
L1	9	350
L1	10	350
E2	9	214
E5A	9	43
E5A	10	43
E5A	11	43
E5B	8	62
E5B	9	62
E5B	10	62
E1	8	402
E4	8	16
E4	10	16
E4	8	9
E4	10	9
E4	11	9
E2	11	168
L1	9	287
E2	8	138
	8	91
E1		26
Ll	8	
L1	9	26
E2	8	131
E2	10	158
E2	11	158
-		

2	3	4		E1	10	612
E5	9	23		E1	9	215
E5	11	23		E1	10	215
E1	10	377		E2	10	298
E1	11	392		E5	8	27
L2	9	237		E5	9	27
L2	8	274		E5	10	27
L2	9	215		E5	11	27
L2	10	215		E2	8	35
E1	9	251		E2	9	35
E1	11	251		E6	8	67
E5	9	22		Еб	11	67
E5	10	22		E6	9	137
E5	11	22		E2	9	295
E5	8	79		E1	8	488
E2	8	72		L1	9	154
E2	11	72		E7 E1	10	71 14
L2	8	285		E1	10	289
L2	11	285		E1	11	289
E1	11	22		E5	. 8	73
E1	11	520		E5	9	73
L1 L2	8	81 417		E5	10	73
	9	417		E6	8	31
L2 E1	10	554		E1	10	640
E1	11	554		E4	9	73
E6	8	37		E6	9	140
E6	10	37		E6	10	140
E4	8	24		E1	8 *	529
E4	9	24		E1	9	529
E4	11	24		E5	8	75
E1	9	319		E5	10	75
L1	11	22		E5	11	75
L2	10	22		E6	8.	104
E1	8 .	525		E6	11	104
E1	9	525		E1	9	385
E6	11	10		E1	10	385
E5	8	11		E1	8	49
E1	10	77		E5	9	39
E1	11	77		E5	11	39
L1	8	349		L2	8	429 429
L1	11	349		L2	9 10	429
E5	9	2.5		L2 E1	8	369
E5	10	25		E1	10	369
E5	11 10	25 101		L1	9	220
E1	9			E2	9	25
L1 E5	8	43 26		L2	10	277
E5	10	26		E6	10	96
E1	10	601		E6	11	96
E1	9	271		E7	8	75
E1	10	271		E7	9	75
E6	11	45		E7	10	75
L1	11	385		E1	10	203
E6	9	47		E1	8	570
L1	8	25		E1	10	570
Ll	9	25		E1	11	570
L1	10	25		L2	9	39 9
E1	9	486		L2	10	346
E1	10	486		E1	11	81

Table X C HPV11 HLA-A24 Supermotif Peptides

			110,1712	Gupermon	r operaco		
E7	9	81			L1	8	444
L1	8	367			E2	10	205
L1	11	367			L2	10	119
E7	10	14			E1	9	339
L1	11	209			E1	8	379
L1	9	439			L1	9	358
E1	11	46			Ll	11	358
Li	10	196			E2	8	130
L2	10	343			E2	9	130
L1	9	199			E2	10	130
E1	9	481			E1	9	613
E1	11	481			E1	11	613
El	8	191			E5	10	67
E1	9	191			E5	11	67
E2	8	96			ь1	9	244
L1	8	332			ь1	10	244
L1	10	332			L1	8	370
E5	8	12			L1	10	370
E5	9	12			E6	8	126
E1	8	534			E1	8	454
L1	11	412			E1	10	454
Li	8	125			E1	11	454
L2	11	29			L2	8	424
L2	11	257			L2	9	424
L2	10	391			E1	9	494
E1	9	236			E5	9	68
L1	9	147			E5	10	68
L1	10	230			E1	10	393
L1	8	365		-	E2	11	345
L1	10	365			El	9	446
E1	9	453			E1	10	446
E1	11	453			E1	8	457
E6	8	113			E1	9	457
E6	9	113			L2	8	238
E1	9	105			L2	11	238
E1	11	105			E5	8	16
E6	10	42			E5	9	20
E2	9	312			E5	11	20
L1	11	454			E1	9	587
E1	8	197			E1	8	220
E6	9	69			L2	8	393
E1	10	604			L2	10	393
E1	11	604			E5	8	23 23
E1	9	131			E5	9	
E2	8	17			E5	10	23
E2	10	17			E5	11	40
El	10	417			E5	8 10	40
E1	11	417			E5		442
E2	9	74			L1	10 9	33
E1	11	360			L1 L1	10	33
E2	11	100			L2	8	430
E2	8	66			L2	9	430
E2	10	66			L1	8	245
E6	10	92			L1	9	245
E6	11	92			L1	10	201
E1	10	128			E1	8	241
E2	9	185			E2	8	360
E1	9	39			L1	8	319
L1	8	103 382			Li	9	319
L1	,	302				-	,,,

E1	10	243	E4	10	36
E1	11	194	L2	8	36
E1	9	326	L2	9	36
E2	9	156	L2	11	36
L1	10	101	L2	11	148
E7	8	22	L2	10	188
E2	8	55	L1	9	362
E1	8	217	L1	10	362
E1	11	217	L1	11	362
L1	10	401	E2	11	183
L2	8	291	E2	11	32
L1	9	145	E1	8	75
L1	11	145	E1	10	412
E1	8	273	L1	9	371
E1	10	273	L1	10	32
L2	8	24	L1	11	32
E1	10	431	E4	11	5
L1	9	408 "	E5	10	34
E1	11	296	L2	9	278
E7	9	85	E1	10	195
L2	11	408.	E6	8	120
E1	8	467	E6	10	120
E1	10	467	L2	8	177
E4	8	99	E5	9	35
E4	10	99	E6	9	97
L1	9	223	E6	10	97
L1	10	223	L2	10	43
L1	11	262	L2	11	43
L2	9	50	E5	8	17
L1	9	111	E5	8	28
L1	10	111	E5	9	28
E1	8	316	E5	10	28
L1	8	113	E1	11	408
L1	11	113	E2	9	30
E1	8	16	L2	8	400
E1	8	44	L2	8	425
L1	11	64	L1	8	377
L2	8	311	L1	10	377
L2	9	311	L1	11	377
E2	10	53	E1	11	56 341
E6	8	119	E1	11	
E6	9	119	L2 L2	9 11	184 184
E6	11	119	L2	10	286
L2	9	176	L2	11	130
E2	8	29	L2	10	188
E2	10	29	E1	10	23
E1	9	264	E5	10	31
E1	11	264	E1	10	443
E2	10	136	E2	11	286
E2	10	78	L2	10	103
E2	10	309	L2	11	103
E5	8	7	L2	9	347
E5	9	7	E6	9	43
E5	10		E2	9	137
E5	11	7 20	L1	10	287
E4	10		E2	8	157
E1	9	305 25	E2	11	157
E6	9	388	L1	9	79
L1			L1	10	79
L1	11	388			

			HLA-A24 Supermon r	epitaes		
E2	10	120		L1	8	303
E1	8	211		E1	11	66
E1	10	211		E5	10	19
E1	8	493		L2	8	158
E1	10	493		L2	8	106
	8	463		E1	8	255
L1		450		E1	11	255
L1	9			E5	8	33
L1	11	450		ES	11	33
E6	8	73		E1	8	557
E6	10	73		E1	10	491
E1	9	312		E5	9	16
E1	10	312				101
E1	11	312		E6	11	
E1	8	254		E5	8	36
E1	9	254		L1	11	187
E6	11	116		Ll	8	41
E6	11	128		L1	9	41
E1	8	357		L1	11	41
E1	10	357		E1	10	521
E1	8	420		E1	9	540
E1	10	420		E2	10	15
E4	8	53		E7	11	83
E6	10	18		E2	8	42
E2	9	84		E2	10	42
E2	11	84		E4	8	18
L1	10	56		E4	9	18
E7	10	39		E1	11	198
E1	8	433		E7	8	82
EI	11	433		E5	9	59
L2	11	33		E5	10	59
E2	9	358		E5	11	59
E2	10	358		E5	11	55
E6	8	99		E5	8	51
L1	10	61		E5	9	51
E2	11	147		E5	10	51
L1	8	19		E5	11	51
Li	10	71		E1	9	298
E6	8	52		E1	10	298
E6	10	52		E1	11	298
L2	8	45		E5	8	69
L2	9	45		E5	9	69
L1	11	122		E5	8	60
E2	8	177		E5	9	60
E2	8	306		E5	10	60
		85		E5	11	60
E1	11	578		E1	9	563
E1	10			E5	10	56
L1	8	227		E5	8	52
E4	8	28		E5	9	52
E1	8	401		E5	10	52
E1	9	401		E5	11	4
E4	8	8		E1	8	514
E4	9	8		E1	9	514
E4	11	8				132
L1	9	87		E1	8	
L1	11	87		E1	9	358
E2	9	310		E5	8	70
E2	11	310		E5	11	70
L1	10	243		E1	9	555
L1	11	243		E1	10	555
E5	9	15		E5	8	49

			 •		
E5	9	49	L1	11	31
E5	10	49	E1	8	300
E5	11	49	E1	9	300
E1	9	268	E2	9	254
E1	10	268	L1	10	446
E2	8	103	E6	10	50
E7	8	48	E1	8	539
L2	8	368	E1	10	539
E6	9	38	E6	9	21
E6	11	38	E2	8	248
E5	8	61	E5	8	54
E5	9	61	E5	9	54
E5	10	61	L2	8	381
L2	9	337	E1	10	97
L2	11	337	L1	11	143
E5	8	47	E2	8	127
E5	9	47	E2	11	127
E5	10	47	E1	9	609
E5	11	47	E1	8	439
L1	8	296	E1	9	439
L1	9	296	E6	9	60
E7	9	5	E6	11	60
E7	11	5	E1	10	594
E1	8	564	E1	8	456
L2	9	245	E1	9	456
E7	10	67	E1	10	456
L1	8	95	L2	9	410
E5	9	57	E4	8	4
E5	11	57	E1	11	442
E1	8	261	E6	8	110
E2	9	18	E6	11	110
E4	8	25	L1	11	218
E4	10	25	L1	9	184
E4	11	25	L2	9	157
E4	9	21	L1	8	459
E4	11	21	L2	10	72
E5	8	53	L1	11	109
E5	9	53	E1	8	562
E4	8	15	E1	10	562
E4	9	15	E5	9	64
E4	11	15	L2	8	385
E1	8	320	L2	9	385
E1	8	306	E1	8	258
E1	10	47	E1	10	258 258
L1	9	197	E1	11 9	513
L1	11	197	E1	10	513
Ll	8	155	E1 E7	8	47
E1	9	274	E7	9.	
E5	8	1	E4	8	68
E1	10	361	L2	10	336
E4 .	9	1	E1	11	545
E4	11	1	L2	9	403
E2	10	101	E2	9	353
L1	10	1	E2	10	353
L2	9	304	L2	11	182
E2	9	170	L1	11	427
E6	11	58	L2	8	123
E1	10	568 451	L2	11	206
E1	8		L1	8	90
E1	11	451		-	

Table X C HPV11 HLA-A24 Supermotif Peptides

			TIEST TE T Daposinom T opinion	
E6	11	87	L2 8 232	
L2	9	95	E1 8 363	
L2	8	170	E6 10 132	
L2	9	170	E1 9 266	
L2	10	170	E1 11 266	
L2	9	422	E2 9 86	
L2	10	422	E6 10 76	
L2	11	422	L1 9 66	
E7	10	20	E6 9 125	
L2	10	358	L2 11 294	
L2	11	358	L2 8 297	
Ll	11	16	L2 9 297	
L2	9	324	L2 8 315	
L2	11	324	E2 9 7	
E5	10	19	E2 10 7	
E5	11	19	E1 8 125	
L2	10	251	L1 11 242	
E2	9	91	E4 11 59	
E2	11	91	L1 9 40	
		80	L1 10 40	
E4	11	238	E2 9 302	
L1	8	238	E2 10 302	
L1	9	238	E1 8 616	
L1	10		E7 8 4	
L1	8	436	E7 10 4	
E5	11	66	L2 10 244	
L1	9	369	E7 11 66	
Ll	11	369	L1 9 94	
E1	10	355	E1 11 324	
E2	9	163	E1 9 293	
E2	8	344	L1 8 473	
E1	8	445	L2 9 220	
E1	10	445	L1 8 141	
E1	11	445	E1 11 583	
L1	8	457	E2 8 300	
Ll	9	457	E2 0 300 E2 11 300	
L1	10	457	L1 8 415	
E5	10	7	E1 9 219	
E5	11	7	L1 11 441	
L2	8	42	E1 9 247	
L2	11	42	E1 9 247 E1 10 247	
E6	11	28	E1 10 247 E2 8 207	
Еб	8	15	E2 8 23	
Еб	10	15		
L1	11	152	E2 9 23 E2 11 23	
L1	8	302	E6 9 12	
Ll	9	302	E6 9 12 E6 11 12	
E2	11	14	E6 11 12 E1 9 547	
E7	10	78	E1 9 547 E1 11 547	
E2	9	288		
E4	8	14		
E4	9	14		
E4	10	14	E1 8 422 L1 9 376	
L1	11	233		
L1	8	251		
E2	9	48		
L1	9	211	E5 11 30	
Ll	11	211	L2 11 102	
E1	8	344	L1 11 286	
L2	9	230	E1 9 350	
L2	10	230	L1 10 86	

			•	•		
L2	9	48		E2	8	354
L2	11	48		E2	9	354
E6	11	23		E2	11	354
L2	8	105		L2	10	183
L2	9	105	-	E5	10	71
E1	11	490		E5	11	71
L2	9	259		L1	8 '	409
L1	9	295		E1	8	294
L1	10	295		E1	10	207
E1	8	260		E1	8	556
E1	9	260		E1	9	556
L2	10	303		E5	8	15
E2	11	260		E.5	10	15
E1	11	206		E7	9	7
E2	9	180		E5	8	50
E2	11	245		E5	9	50
L2	8	209		E5	10	50
L2	10	370		E5	11	50
L2	9	240		E1	10	297
El	11	185		E1	11	297
E6	10	7		E7	8	86
E4	8	85		E2	9	93
E1	10	331		E2	10	93
E1	11	331		E2	11	93
E2	8	201		E6	8	26
E2	8	151		L1	9	378
E2	9	151		L1	10	378
E2	9	257		L1	10	389
E1	8	60		E6	8	39
L2	8	152		E6	10	39
E1	8	436		E1	11	232
E1	11	436		E6	8	142
L1	8	190		E1	9	585
E2	8	348		E1	11	585
L1	8	393		L2	8	37
L1	10	3 9 3		L2	10	37
E2	10	40		E5	9	14
E5	8	45		E5	11	14
E5	9	45		E5	8.	62
E5	10	45		E5	9	62
E5	11	45		E5	11	62
Ll	10	176		E1	11	116
L1	8	344		L1	9	333
E2	11	198		L2 L2	10	150 150
L2	11	144		E5	10	130
L2	9	378		L2	10	149
L2 L2	11 10	378		L2	11	149
		366		E5	11	12
E1	9	501		E5	8	55
E1	10	501		L2	11	382
E1	1 1 9	501 344		E4	9	100
L2				E1	9	290
E6	9	19		E1	10	290
E6	11	19		L1	8	224
E1 L2	8	588 401		L1	9	224
	11			L1	11	224
E1 E1	8 10	586 586		E6	10	88
L2	9	38		E6	11	88
E2	10	261		L1	10	263
E-2	10	201				

					-	
Ll	11	263		L2	11	414
E1	9	332		L2	8	325
E1	10	332		L2	10	325
L2	10	383		E2	10	148
L2	11	383		E2	11	148
E1	9	78		E1	10	584
El	10	78		L2	8	246
E4	9	82		E1	11	238
E4	11	32		E5	8	24
E2	11	63		E5	9	24
E4	8	101		E5	10	24 330
L1	11	329		L1 E7	10 9	68
L1	9	57		E5	9	20
L1	11	57			10	20
E5	8	13		E5 L1	9	72
E5	11	13		L1	11	72
E1	10	531		E4	8	2
E1	11	531		E4	10	2
L1	10	413		L1	8	58
E2	11	335		Li	10	58
L1	10	350		L2	9	120
L1	11	350		L2	11	120
L2	8	51		E6	9	53
L2	8	423		E2	8	132
L2	9	423		E5	9	41
L2	10	423		E5	11	41
E5	8	21 21		E4	8	74
E5	9			E6	8	54
E5	11	21 346		L2	9	252
E1	11	333		L2	11	252
E1	8 9	333		E1	9	602
E1 L2	10	30		E5	11	60
E1	11	499		E5	8	42
L1	9	394		E5	10	42
E5	9	27		E5 -	11	42
E5	9	32		L2	9	392
E2 .	9	41		L2	11	392
E2	11	41		E1	8	340
E4	9	17		E5	10	14
E4	10	17		E5	11	18
E1	8	275		E1	11	464
Li	8	73		E5	8	58
Li	10	73		E5	10	58
E5	8	48		E5	11	58
E5	9	48		E1	8	267
E5	10	48		E1	10	267
E5	11	48		E1	11	267
E5	8	46		E2	9	102
E5	9	46		E2	8	92
E5	10	46		E2	10	92
E5	11	46		E2	11	92
E4	9	10		E6	8	141
E4	10	10		E6	9	141
Ll	8	383		E4	10	81
E2	10	128		E1	8	530
E2	11	128		E1	11	530
E7	8	6		E1	8	237
E7	10	6		L2 .	11	436
L2	10	145		L2	10	437

E5	9		76
E5	10		76
E5	11		76
L1	10		386
E4	ė.		22
E4	10		22
E4	11		22
E6	10		105
E1	10		86
L2	8		431
E1	9		579
E1	11		579
E5	8	•	54
E2	8		138
L1	9		231
L1	8		246
L2	9		367
E1	9		532
E1	10		532
Ll	8		359
L1	10		359
E1	11		536
E5	10		61
E2	9		159
E2	10		159
L1	9		351
L1	10		351
L2	8		305
E5	9		43
E5	10		43
E5	11		43
Ll	9		288
E1	8		402
Ll	8		26
Ll	9		26
E4	8		16
E4	10		16
E4	11		16
E4	8		9
E4	10		9
E4	11		9
E2	11		168
E1	8		502 502
E1	9		
E1	10 8		502 131
E2	9		131
E2 E2			158
E2 E2	10		158
EZ	11		128

Table XI HLA-B7 Supermotif Peptides

				WD111 6 D2	8	5
1	2	3	4	HPV16 E7 HPV16 E7	9	5
HPV16		10	559 308	HPV16 E7	11	5
HPV16		9	308	HPV16 L1	10	461
HPV16		10	308	HPV16 L1	10	211
HPV16		11	592	HPV16 L1	11	211
HPV16		10	592	HPV16 L1	-8	465
HPV16		11	467	HPV16 L1	9	465
HPV16		9	467	HPV16 L1	11	465
HPV16		10	467	HPV16 L1	8	266
HPV16		11	599	HPV16 L1	9	266
HPV16		8	309	HPV16 L1	10	266
HPV16		9	309	HPV16 L1	11	266
HPV16		10	309	HPV16 L1	11	76
HPV16		11	309	HPV16 L1	8	488
HPV16		9	560	HPV16 L1	9	488
HPV16		11	560	HPV16 L1	11	488
HPV16		8	511	HPV16 L1	8	318
HPV16		10	511	HPV16 L1	9	318
HPV16		11	511	HPV16 L1	11	503
HPV16		11	552	HPV16 L1	8	80
HPV16		10	93	HPV16 L1	9	80
HPV16		9	107	HPV16 L1	8	188
HPV16	E1	10	336	HPV16 L1	8	335
HPV16	E1	11	336	HPV16 L1	9	103
HPV16	E1	9	189	HPV16 L1	9	39
HPV16	E1	10	189	HPV16 L1	10	39
HPV16	E1	8	244	HPV16 L1	8	31
HPV16	E1	11	244	HPV16 L1	9	31
HPV16	E1	11	526	HPV16 L1	8	118
HPV16	E1	8	576	HPV16 L1	9	118
HPV16	E1	9	576	HPV16 L1	10	118
HPV16		10	576	HPV16 L1	8 10	207
HPV16		11	576	HPV16 L1 HPV16 L1	11	207
HPV16		11	105	HPV16 L1	9	459
HPV16		11	195	HPV16 L1	10	435
HPV16		9	218	HPV16 L1	9	212
HPV16		10	218	HPV16 L1	10	212
HPV16		11	218	HPV16 L1	11	434
HPV16		11	352	HPV16 L1	8	40
HPV16		8	249 249	HPV16 L1	9	40
HPV16		9	249	HPV16 L1	8 .	433
HPV16 HPV16		9 10	207	HPV16 L1	8	199
HPV16		10	286	HPV16 L1	10	458
HPV16		11	59	HPV16 L1	11	320
HPV16		10	69	HPV16 L1	11	514
HPV16		10	30	HPV16 L1	9	88
HPV16		9	118	HPV16 L1	11	88
HPV16		9	11	HPV16 L1	9	246
HPV16		8	101	HPV16 L1	8	257
HPV16		11	101	HPV16 L2	9	278
HPV16		8	19	HPV16 L2	10	278
HPV16		10	65	HPV16 L2	11	278
HPV16		8	15	HPV16 L2	8	424
HPV16		9	46	HPV16 L2	8	119
HPV16		10	46	HPV16 L2	9	87
HPV16		11	40	HPV16 L2	9	28
HPV16		8	16	HPV16 L2	11	239
HPV16	E7	10	16	HPV16 L2	8	280

Table XI HLA-B7 Supermotif Peptides

HPV16	L2	9	280		HPV16 L			123
HPV16	L2	10	280		HPV16 I			123
HPV16	L2	8	102		HPV16 I			385
HPV16	L2	11	165		HPV16 I			385
HPV16	L2	10	96		HPV16 I			382
HPV16		11	96		HPV18 F	1		566
HPV16		8	413		HPV18 E	1	8	518
HPV16		8	99		HPV18 B	31	9	518
HPV16		10	99		HPV18 B	31	10	518
HPV16		11	99		HPV18 E	1	11	518
HPV16		9	394		HPV18 E	31	10	599
HPV16		9	400		HPV18 E	81	11	599
HPV16		11	400		HPV18 E	31	9	484
HPV16		8	216			51	10	484
HPV16		10	216			21	8	576
		9	416			31	10	576
HPV16		10	428			31	11	576
HPV16			33			31	11	559
HPV16		9				31	8	641
HPV16		10	33		HPV18 H		10	641
	L2	10	73			31	9	193
	L2	9	408			31	8	251
HPV16		11	408			51 51	11	251
HPV16		8	196			51 51	11	606
HPV16		8	126			31	9	567
HPV16	L2	10	126					316
HPV16	L2	9	357			31	8	316
HPV16	L2	10	357			31	9	316
HPV16	L2	8	462			31		
HPV16	L2	10	462	•		31	11	316
HPV16	L2	11	462			31	9	263
HPV16	L2	9	254			E1	9	315
HPV16	L2	11	206			E1	10	315
HPV16	L2	9	160			E1	11	315
HPV16	L2	10	160			E1	9	447
HPV16	L2	8	29			B1	9	97
HPV16	L2	9	127			B1	9	110
HPV16	L2	8	91			E1	8	343
HPV16		8	79			E1	10	343
HPV16		9	79			B1	11	343
HPV16	L2	11	79		HPV18		8	474
HPV16		11	171			E1	9	474
HPV16		11	291		HPV18	B1	8	307
HPV16		9	90			E1	8	583
HPV16		9	78		HPV18	E1	10	583
HPV16		10	78		HPV18	E1	11	583
HPV16		8	220		HPV18	E2	9	261
	L2	9	220		HPV18	E2	10	261
HPV16		8	434		HPV18	E2	8	352
HPV16		10	434		HPV18	E2	9	352
HPV16		9	173		HPV18	E2	11	352
HPV16		11	173		HPV18	E2	8	248
HPV16		11	142		HPV18	E2	9	226
	L2	9	214			E2	10	3
HPV16		10	214			E2	9	224
		11	345			E2	11	224
	L2	8	245			E2	11	271
HPV16	L2		380			E2	11	63
	L2	11	431			E5	8	59
	L2	9	431			E5	10	59
HPV16		10	431		HPV18		11	59
HPV16	L 2	11	#31				-	

Table XI HLA-B7 Supermotif Peptides

				•			
HPV18	E5	8	23	HPV18	L1		343
HPV18	E5	9	23	HPV18			153
HPV18	E5	10	23	HPV1'8		9	153
HPV18	E5	11	23	HPV18			153
HPV18		8	45	HPV18		8	108
HPV18	E5	9	45	HPV18		9	247
HPV18		10	45	HPV18		10	247
HPV18	E5	11	45	HPV18		8	469
HPV18		9	20	HPV18		9	469 75
HPV18		11	20	HPV18		9	75
HPV18		9	6	HPV18		8	470
HPV18		10	60	HPV18		11	470
HPV18		9	110	HPV18		8	76
HPV18		8	14	HPV18		10	471
HPV18		9	113	HPV18		8	25
HPV18		8	10	HPV18		9	25
HPV18		8	16	HPV18		10	25
HPV18		10	16 -	HPV18		11	25
HPV18		11	16	HPV18		11	239
HPV18		10	5 5 5 5	HPV18		8	66
HPV18		9	3	HPV18		9	66
HPV18 HPV18		10	3	HPV18		8	353
HPV18		11	3	HPV18		9	353
HPV18		11	21	HPV18		9	411
HPV18		9	495	HPV18		8	398
HPV18		8	223	HPV18		10	398
HPV18		9	223	HPV18	L1	9	90
HPV18		11	549	HPV18	L1	11	113
HPV18		10	246	HPV18	L1	10	413
HPV18		11	246	HPV18	L1	9	281
HPV18		8	501	HPV18	L1	9	468
HPV18		9	501	HPV18	L1	10	468
HPV18	L1	11	501	HPV18	L1	9	292
HPV18		8	301	HPV18		8	524
HPV18	L1	9	301	HPV18		9	524
HPV18		11	301	HPV18		11	524
HPV18	L1	8	56	HPV18		8	421
HPV18	L1	9	56	HPV18		10	421
HPV18	L1	10	56	HPV18		9	327
HPV18	L1	9	19	HPV18		9	27
HPV18	L1	11	19	HPV18		11	266 100
HPV18		10	543	HPV18		8	208
HPV18		10	23	HPV18 HPV18		11	208
HPV18		11	23	HPV18		9	257
HPV18		9	123	HPV18		10	94
HPV18		11	123	HPV18		11	94
HPV18		8	557	HPV18		8	223
HPV18		10	557	HPV18		9	223
HPV18		11	539	HPV18		8	97
HPV18		8	370	HPV18		10	97
HPV18		9	138 27	HPV18		11	97
HPV18		8		HPV18		8	85
HPV18		9	27 27	HPV18		9	85
HPV18		10	27	HPV18		8	444
HPV18		11		HPV18		10	444
HPV18		8 9	15 74	HPV18		11	444
HPV18			74	HPV18		10	72
HPV18		10 9	343	HPV18		11	72
HPV18	11	9	343		-		

Table XI HLA-B7 Supermotif Peptides

HPV18	L2	8	407	HPV3		10	491
HPV18		10	407 .	HPV3:		11	491
HPV18		11	407	HPV3		9	288
HPV18	L2	8	215	HPV3		10	288
HPV18	L2	11	253	HPV3		11	288
HPV18	L2	9	159	HPV3		9	236
HPV18	L2	10	159	HPV3		10	92
HPV18	L2	11	238	HPV3		9	106
HPV18	L2	8	28	HPV3		10	506
HPV18	L2	8	89	HPV3		11	506
HPV18	L2	11	284	HPV3		10	316 316
HPV18	L2	9	88	HPV3		11 9	169
HPV18		8	244	HPV3		10	169
HPV18		10	119	HPV3		8	447
HPV18		10	329	HPV3 HPV3		9	447
HPV18		11	329	HPV3		8	556
HPV18		9	324	HPV3		9	556
HPV18		10	324	HPV3		10	556
HPV18		10	418	HPV3		11	556
HPV18		11	418	HPV3		11	105
HPV18		10	375	HPV3		8	289
HPV18		9	213	HPV3		10	289
HPV18		10	213	HPV3		11	195
HPV18		9	144	HPV3		11	257
HPV18		9	184	HPV3		11	359
HPV18		9	271 271	HPV3		10	293
HPV18		10	32	HPV3		9	59
HPV18		9 10	32	HPV3		11	59
HPV18 HPV18		9	389	HPV3		10	69
HPV18		11	389	HPV3	1 E5	9	30
HPV18		8	170	HPV3	1 E5	10	30
HPV18		11	170	HPV3	1 E5	11	30
HPV18		9	361	HPV3	1 E5	8	55
HPV18		10	361	HPV3	1 E5	9	55
HPV18		11	361	HPV3	1 E5	10	55
HPV18		11	359	HPV3	1 E5	11	55
HPV18		9	397	HPV3	1 E6	9	111
HPV18		11	397	HPV3		8	21
HPV18		9	451	HPV3		11	21
HPV18		10	451	HPV3		9	4
HPV18	L2	11	451	HPV3		8	8
HPV18	L2	8	414	HPV3		11	8
HPV18	L2	10	414	HPV3		8	142 58
HPV31	El	8	458	HPV3		10 8	91
HPV31		9	458	HPV3		9	46
HPV31	E1	11	458		1 E7	10	46
HPV31		10	539	HPV3		10	28
HPV31		10	572		1 E7	11	28
HPV31		11	572	HPV3		10	16
HPV31		11	128	HPV3		8	5
HPV31		11	523	HPV3		9	5
HPV31		11	579	HPV3		11	5
HPV31		8	289	HPV		10	433
HPV31		9	289		31 L1	10	488
HPV3 1		10	289		31 L1	10	186
HPV31		11	289 540		31 L1	11	186
HPV31		9			31 L1	8	440
HPV31		11	540		31 L1	9	440
. HPV31	E1	8	491			-	

Table XI HLA-B7 Supermotif Peptides

HPV31	L1	11	440	HPV31		11	234
HPV31	L1	8	241	HPV31		10	96
HPV31	L1	9	241	HPV31		11	96
HPV31	L1	10	241	HPV31		10	421
HPV31		11	241	HPV31		9	406
HPV31		8	463	HPV31		8	99
HPV31	L1	9	463	HPV31		10	99 99
HPV31		11	463	HPV31		11	125
HPV31		8	293	HPV31		8	125
HPV31		9	293	HPV31 HPV31		10 11	125
HPV31		10	139	HPV31		8	211
HPV31		10	52	HPV31		10	211
HPV31		11	52	HPV31		10	123
HPV31		10	436	HPV31		9	401
HPV31		11	436	HPV31		11	401
HPV31		8	163	HPV31		9	87
HPV31		8	310	HPV31		9	33
HPV31		9	78	HPV31		10	33
HPV31		9	13	HPV31		8	191
HPV31		10	13	HPV31		9	327
HPV31		8	396	HPV31		9	249
HPV31		10	396	HPV31		9	155
HPV31		8	93	HPV31		10	155
HPV31		9	93	HPV31		11	166
HPV31		10	93 57	HPV31		9	126
HPV31		10		HPV31		10	126
HPV31		9	187	HPV31		8	91
HPV31		10	187 410	HPV31		11	91
HPV31		10		HPV31		11	284
HPV31		8	14 14	HPV31		8	216
HPV31		9	478	HPV31		11	216
HPV31		11	5	HPV31		9	90
HPV31		9	5	HPV31		8	78
HPV31		8	174	HPV31		10	78
HPV31 HPV31		8	182	HPV31		11	372
HPV31		10	182	HPV31		9	168
HPV31		11	409	HPV31	L2	10	168
HPV31		11	295	HPV31	L2	11	168
HPV31		9	63	HPV31	L2	10	141
HPV31		11	63	HPV31	L2	10	209
HPV31		9	221	HPV31	L2	8	427
HPV31		8	232	HPV31	L2	10	409
HPV31		9	232	HPV31	L2	11	409
HPV31		9	271	HPV31	L2	9	393
HPV31		10	271	HPV31	L2	11	393
HPV31		11	271	HPV31		10	73
HPV31	L2	8	240	HPV31		9	387
HPV31	L2	8	414	HPV33		8	3
HPV31	L2	10	414	HPV33		10	3
HPV31	L2	11	414	HPV33		11	301
HPV31	L2	9	424	HPV33		10	585
HPV31		10	424	HPV33		11	585
HPV31		11	424	HPV33		9	470
HPV31		9	28	HPV33		10	470
HPV31	L2	8	273	HPV33		8	293
HPV31	L2	9	273	HPV33		8	460
HPV31	L2	10	273	HPV33		9	460
HPV31		8	102	HPV33		11	592
HPV31	L2	11	160	HPV33	ы	10	302

Table XI HLA-B7 Supermotif Peptides

HPV33	E1	11	302	HPV33 L1	11	180
HPV33	E1	11	553	HPV33 L1	10 -	483
HPV33	E1	8	504	HPV33 L1	10	185
HPV33	E1	10	504	HPV33 L1	11	185
HPV33		11	504	HPV33 L1	8	438
HPV33		9	433	HPV33 L1	9	438
HPV33		8	237	HPV33 L1	11	438
HPV33		10	237	HPV33 L1	8	240
HPV33		11	237	HPV33 L1	9	240
HPV33		8	329	HPV33 L1	10	240
HPV33		10	329	HPV33 L1	11	240
HPV33		11	329	HPV33 L1	8	461
HPV33		8	518	HPV33 L1	9	461
		11	518	HPV33 L1	11	461
HPV33			569	HPV33 L1	8	292
HPV33		8	569	HPV33 L1	9	292
HPV33		10		HPV33 L1	8	476
HPV33		11	569	HPV33 L1	8	163
HPV33		10	195	HPV33 L1	8	309
HPV33		9	247	HPV33 L1	9	78
HPV33		11	340		9	13
HPV33		10	294	HPV33 L1		13
HPV33		8	26	HPV33 L1	10	394
HPV33	E2	10	26	HPV33 L1	8	
HPV33	E2	11	26	HPV33 L1	10	394
HPV33	E2	10	341	HPV33 L1	8	93
HPV33	E2	11	341	HPV33 L1	9	93
HPV33	E2	9	238	HPV33 L1	10	93
HPV33	E2	9	229	HPV33 L1	8	54
HPV33		11	59	HPV33 L1	9	54
HPV33	E5	8	59	HPV33 L1	10	54
HPV33		10	59	HPV33 L1	9	432
HPV33		10	20	HPV33 L1	9	186
HPV33		11	20	HPV33 L1	10	186
HPV33		9	45	HPV33 L1	11	407
HPV33		10	45	HPV33 L1	10	408
HPV33		11	45	HPV33 L1	11	164
HPV33		9	111	HPV33 L1	8	14
HPV33		11	111	HPV33 L1	9	14
HPV33		8	94	HPV33 L1	10	139
HPV33		11	94	HPV33 L1	8	5
		8	35	HPV33 L1	9	5
HPV33		9	35	HPV33 L1	8	406
HPV33		11	35	HPV33 L1	9	63
HPV33			8	HPV33 L1	11	63
HPV33		8	8	HPV33 L1	10	431
HPV33		11	58	HPV33 L2	9	173
HPV33		8		HPV33 L2	10	173
HPV33		10	58	HPV33 L2	11	173
HPV33		8	18	HPV33 L2	9	276
HPV33		11	18	HPV33 L2	10	276
HPV33		8	5	HPV33 L2	11	276
HPV33		9	5			120
HPV33		11	5	HPV33 L2	10	27
HPV33		8	46	HPV33 L2	9	239
HPV33	E7	9	46	HPV33 L2	11	
HPV33	E7	10	46	HPV33 L2	8	278
HPV33	E7	9	40	HPV33 L2	9	278
HPV33	E7	11	40	HPV33 L2	10	278
HPV33	E7	8	16	HPV33 L2	8	261
HPV33	E7	10	16	HPV33 L2	9	77
HPV33		9	180	HPV33 L2	9	165
155		-				

PCT/US00/33549

Table XI HLA-B7 Supermotif Peptides

HPV33 L2					nlA-b/ S	upermont rep	lides			
HPV33 L2	HPV33	T.2	11	165		н	PV33	L2	11	338
HPV33 L2						Н	PV45	E1	10	552
HPV33 L2			11	428						
HPV33 L2	HPV33	L2	8	415						
New			10	415						
New	HPV33	L2	8	456		H	PV45	E1		
NPV33 L2	HPV33	L2	10	456						
NPV33 L2	HPV33	L2	11	456						
NPV33 L2	HPV33	L2	8	407						
HPV33 L2	HPV33	L2	9	407						
NPV33 L2	HPV33	L2	8	98						
HPV33 L2	HPV33	L2	10							
NPV33 L2	HPV33	L2	11							
HPV43 L2	HPV33	L2								
HPV33 L2										
NPV33 L2										
NPV33 L2										
HPV43 L2										
HPV33 L2										
HPV33 L2										
HPV33 L2 9 160 HPV45 E1 11 592 HPV33 L2 10 160 HPV45 E1 1 11 592 HPV33 L2 11 171 HPV45 E1 8 302 HPV33 L2 11 171 HPV45 E1 1 0 302 HPV33 L2 8 78 HPV45 E1 10 302 HPV33 L2 11 78 HPV45 E1 11 302 HPV33 L2 8 90 HPV45 E1 11 302 HPV33 L2 11 90 HPV45 E1 9 433 HPV33 L2 9 86 HPV45 E1 9 110 HPV33 L2 11 289 HPV45 E1 9 110 HPV33 L2 11 289 HPV45 E1 8 329 HPV33 L2 11 289 HPV45 E1 8 329 HPV33 L2 11 289 HPV45 E1 10 329 HPV33 L2 11 289 HPV45 E1 10 329 HPV33 L2 11 289 HPV45 E1 8 329 HPV33 L2 11 289 HPV45 E1 8 329 HPV33 L2 9 89 HPV45 E1 11 329 HPV33 L2 9 E20 HPV45 E1 8 460 HPV33 L2 9 E20 HPV45 E1 8 293 HPV33 L2 9 E20 HPV45 E1 8 293 HPV33 L2 9 E24 HPV45 E1 11 569 HPV33 L2 11 274 HPV45 E1 8 569 HPV33 L2 9 E25 HPV45 E2 11 569 HPV33 L2 9 E25 HPV45 E2 11 569 HPV33 L2 11 425 HPV45 E2 11 569 HPV33 L2 10 425 HPV45 E2 11 569 HPV33 L2 10 429 HPV45 E2 11 569 HPV33 L2 10 419 HPV45 E2 10 66 HPV33 L2 10 419 HPV45 E2 10 569 HPV33 L2 10 419 HPV45 E2 10 58 H										
HPV33 L2										
HPV33 L2										
HPV33 L2										
HPV33 L2										
HPV33 L2										
HPV33 L2 8 90 HPV45 E1 9 433 HPV33 L2 11 90 HPV45 E1 9 97 HPV33 L2 9 86 HPV45 E1 9 91 HPV33 L2 11 346 HPV45 E1 9 110 HPV33 L2 11 289 HPV45 E1 10 329 HPV33 L2 8 220 HPV45 E1 11 329 HPV33 L2 8 220 HPV45 E1 11 329 HPV33 L2 8 220 HPV45 E1 10 460 HPV33 L2 9 220 HPV45 E1 11 329 HPV33 L2 8 274 HPV45 E1 10 460 HPV33 L2 8 274 HPV45 E1 8 293 HPV33 L2 10 274 HPV45 E1 8 293 HPV33 L2 9 274 HPV45 E1 8 293 HPV33 L2 9 1274 HPV45 E1 10 569 HPV33 L2 11 274 HPV45 E1 11 569 HPV33 L2 10 425 HPV45 E1 11 569 HPV33 L2 10 425 HPV45 E1 11 569 HPV33 L2 10 425 HPV45 E2 11 5569 HPV33 L2 10 425 HPV45 E2 11 5569 HPV33 L2 10 425 HPV45 E2 11 5569 HPV33 L2 10 419 HPV45 E2 8 240 HPV33 L2 10 419 HPV45 E2 10 65 HPV33 L2 10 419 HPV45 E2 10 55 HPV33 L2 10 419 HPV45 E2 10 65 HPV33 L2 10 412 HPV45 E2 10 55 HPV33 L2 10 412 HPV45 E2 10 65 HPV33 L2 10 412 HPV45 E2 11 65 HPV33 L2 10 138 HPV45 E6 9 6 HPV33 L2 10 138 HPV45 E6 9 6 HPV33 L2 10 138 HPV45 E6 9 6 HPV33 L2 10 138 HPV45 E7 8 22 HPV33 L2 10 175 HPV45 E7 8 22 HPV33 L2 10 175 HPV45 E7 9 16 HPV33 L2 10 175 HPV45 E7 9 16 HPV33 L2 10 72 HPV45 E7 9 16 HPV33 L2 9 422 HPV45 E7 9 3										
HPV33 L2										
HPV33 LZ 9 866 HPV45 E1 8 129 HPV33 LZ 11 346 HPV45 E1 10 329 HPV33 LZ 11 289 HPV45 E1 11 329 HPV33 LZ 12 8 220 HPV45 E1 8 460 HPV33 LZ 8 220 HPV45 E1 9 460 HPV33 LZ 8 274 HPV45 E1 8 569 HPV33 LZ 8 274 HPV45 E1 8 569 HPV33 LZ 11 274 HPV45 E1 8 569 HPV33 LZ 9 274 HPV45 E1 8 569 HPV33 LZ 9 425 HPV45 E1 8 569 HPV33 LZ 11 274 HPV45 E1 10 569 HPV33 LZ 9 425 HPV45 E1 11 569 HPV33 LZ 9 425 HPV45 E1 11 569 HPV33 LZ 10 425 HPV45 E1 11 569 HPV33 LZ 10 425 HPV45 E2 9 355 HPV33 LZ 10 425 HPV45 E2 9 355 HPV33 LZ 10 425 HPV45 E2 10 569 HPV33 LZ 10 429 HPV45 E2 10 56 HPV33 LZ 10 429 HPV45 E2 10 56 HPV33 LZ 10 329 HPV45 E2 10 66 HPV33 LZ 10 412 HPV45 E2 10 56 HPV33 LZ 10 412 HPV45 E2 10 66 HPV33 LZ 11 412 HPV45 E2 10 68 HPV33 LZ 10 412 HPV45 E2 10 238 HPV33 LZ 10 412 HPV45 E2 11 65 HPV33 LZ 10 412 HPV45 E2 11 65 HPV33 LZ 10 138 HPV45 E6 9 6 HPV33 LZ 10 138 HPV45 E6 9 6 HPV33 LZ 10 138 HPV45 E6 9 113 HPV33 LZ 10 214 HPV45 E6 9 113 HPV33 LZ 10 214 HPV45 E7 8 22 HPV33 LZ 10 175 HPV45 E7 8 22 HPV33 LZ 10 175 HPV45 E7 9 16 HPV33 LZ 11 125 HPV45 E7 9 16 HPV33 LZ 11 125 HPV45 E7 9 16 HPV33 LZ 11 125 HPV45 E7 9 16 HPV33 LZ 10 72 HPV45 E7 10 56 HPV35 LZ 11 125 HPV45 E7 9 16 HPV33 LZ 10 72 HPV45 E7 10 56 HPV35 LZ 10 72 HPV45 E7 10 58										
HPV33 L2										110
HPV35 L2									8	329
HPV35 L2									10	329
HPV33 L2								E1	11 .	329
HPV33 L2						F	IPV45	E1	8	460
HPV33 L2						F	IPV45	E1	9	460
## ## ## ## ## ## ## ## ## ## ## ## ##						I	IPV45	E1	10	460
HPV33 L2								E1	8	293
HPV33 L2						1	IPV45	E1	8	
HPV33 L2						I	IPV45	E1		
HPW33 L2 11 425 HPW45 E2 9 355 HPW45 E2 11 1555 HPW35 L2 10 419 HPW45 E2 11 1555 HPW35 L2 10 419 HPW45 E2 10 66 HPW35 L2 10 419 HPW45 E2 10 66 HPW35 L2 10 412 HPW45 E2 10 5 HPW35 L2 10 412 HPW45 E2 10 238 HPW35 L2 11 412 HPW45 E2 11 65 HPW33 L2 11 412 HPW45 E2 11 65 HPW33 L2 10 138 HPW45 E2 11 65 HPW33 L2 10 138 HPW45 E6 9 16 HPW33 L2 10 138 HPW45 E6 8 14 HPW33 L2 10 214 HPW45 E6 8 114 HPW33 L2 10 214 HPW45 E6 8 10 HPW33 L2 10 214 HPW45 E7 8 22 HPW33 L2 10 375 HPW45 E7 8 22 HPW33 L2 11 375 HPW45 E7 8 22 HPW33 L2 11 375 HPW45 E7 8 22 HPW33 L2 11 125 HPW45 E7 9 16 HPW35 L2 11 125 HPW45 E7 9 16 HPW35 L2 11 72 HPW45 E7 8 56 HPW45 E7 11 12 HPW33 L2 10 72 HPW45 E7 11 65 HPW35 L2 11 72 HPW45 E7 8 56 HPW45 E7 11 67 HPW35 L2 11 72 HPW45 E7 8 56 HPW45 E7 11 16 HPW35 L2 11 72 HPW45 E7 8 56 HPW45 E7 10 56 6 HPW35 L2 11 72 HPW45 E7 8 56 HPW35 L2 11 72 HPW45 E7 9 3 3 HPW35 L2 9 422 HPW45 E7 9 3 3 10 10 10 10 10 10 10 10 10 10 10 10 10				425		F	1PV45			
HPV33 L2			11	425		1	IPV45			
HPV33 L2			9	419						
HPW33 L2 8 245 HPW45 E2 10 66 HPW33 L2 10 329 HPW45 E2 10 5 HPW33 L2 11 412 HPW45 E2 11 23 HPW33 L2 11 412 HPW45 E2 11 65 HPW33 L2 9 1.85 HPW45 E6 9 6 HPW33 L2 10 13.8 HPW45 E6 9 6 HPW33 L2 10 13.8 HPW45 E6 9 11 HPW33 L2 10 214 HPW45 E6 9 113 HPW33 L2 10 214 HPW45 E6 8 10 HPW33 L2 10 375 HPW45 E7 8 22 HPW33 L2 11 375 HPW45 E7 8 22 HPW33 L2 11 125 HPW45 E7 9 16 HPW33 L2 11 172 HPW45 E7 8 56 HPW33 L2 10 72 HPW45 E7 1 16 HPW33 L2 10 72 HPW45 E7 10 56 HPW33 L2 10 72 HPW45 E7 10 56 HPW33 L2 11 72 HPW45 E7 8 56			10	419						
HPV33 L2 10 329 HPV45 E2 10 5 HPV33 L2 10 412 HPV45 E2 11 65 HPV33 L2 11 412 HPV45 E2 11 65 HPV33 L2 10 138 HPV45 E6 9 6 HPV33 L2 10 138 HPV45 E6 9 113 HPV33 L2 10 214 HPV45 E6 8 14 HPV33 L2 10 214 HPV45 E6 8 10 HPV33 L2 10 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 11 12 HPV33 L2 11 125 HPV45 E7 11 12 HPV33 L2 11 125 HPV45 E7 11 16 HPV33 L2 11 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 9 3			8	245						
HPV33 L2 10 412 HPV45 E2 11 65 HPV33 L2 9 185 HPV45 E6 9 6 HPV45 E6 11 65 HPV45 E2 11 65 HPV33 L2 10 138 HPV45 E6 9 14 HPV45 E6 9 11 HPV33 L2 10 214 HPV45 E6 9 113 HPV33 L2 10 214 HPV45 E7 8 10 HPV33 L2 10 275 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 11 22 HPV45 E7 11 12 HPV33 L2 11 375 HPV45 E7 9 16 HPV33 L2 11 375 HPV45 E7 9 16 HPV33 L2 11 375 HPV45 E7 9 16 HPV33 L2 11 372 HPV45 E7 11 16 HPV33 L2 11 372 HPV45 E7 11 16 HPV33 L2 11 372 HPV45 E7 11 16 HPV33 L2 11 72 HPV45 E7 10 56 HPV35 L2 11 72 HPV45 E7 10 56 HPV35 L2 9 422 HPV45 E7 9 3 HPV35 L2 9 422 HPV45 E7 9 3			10	329						
HPV33 L2 11 412 HPV45 E6 9 6 HPV33 L2 10 138 HPV45 E6 8 14 HPV33 L2 9 214 HPV45 E6 9 113 HPV33 L2 10 214 HPV45 E6 8 10 HPV33 L2 10 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 11 125 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 1 56 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 1 56 HPV33 L2 11 72 HPV45 E7 1 56 HPV33 L2 9 422 HPV45 E7 9 56	HPV33	L2	10	412						
HPV33 L2 9 105 HPV45 E6 8 14 HPV33 L2 9 214 HPV45 E6 8 10 HPV33 L2 10 214 HPV45 E6 8 110 HPV33 L2 10 214 HPV45 E6 8 10 HPV33 L2 10 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 8 12 HPV33 L2 11 375 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 9 3	HPV33	L2	11	412						
HPV33 L2 10 136 HPV45 E6 9 113 HPV33 L2 10 214 HPV45 E6 8 10 HPV33 L2 10 375 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 11 16 HPV33 L2 9 402 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 8 56	HPV33	L2	9	185						
HPV33 L2 9 214 HPV45 E6 8 10 HPV33 L2 10 214 HPV45 E7 8 22 HPV33 L2 11 375 HPV45 E7 11 22 HPV33 L2 11 375 HPV45 E7 9 16 HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 9 402 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 9 3	HPV33	L2	10							
HPV33 LZ 10 375 HPV45 E7 8 22 HPV33 LZ 11 375 HPV45 E7 11 22 HPV33 LZ 11 125 HPV45 E7 9 16 HPV33 LZ 9 402 HPV45 E7 11 16 HPV33 LZ 9 402 HPV45 E7 11 16 HPV33 LZ 10 72 HPV45 E7 8 56 HPV33 LZ 11 72 HPV45 E7 10 56 HPV33 LZ 9 422 HPV45 E7 9 3	HPV33	L2	9	214						
HPV33 L2 10 375 HPV45 E7 11 22 HPV45 L2 11 375 HPV45 E7 11 22 HPV33 L2 11 125 HPV45 E7 11 16 HPV33 L2 9 402 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 9 3 HPV35 L2 9 422 HPV45 E7 9 3										
HPV33 L2 11 125 HPV45 E7 9 16 HPV33 L2 9 402 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 9 422 HPV45 E7 9 3										
HPV33 L2 9 402 HPV45 E7 11 16 HPV33 L2 10 72 HPV45 E7 8 56 HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 11 72 HPV45 E7 9 3 HPV3 L2 9 422 HPV45 E7 9 3	HPV33	L2								
HPV33 L2 10 72 HPV45 B7 8 56 HPV33 L2 11 72 HPV45 B7 10 56 HPV33 L2 9 422 HPV45 B7 9 3										
HPV33 L2 11 72 HPV45 E7 10 56 HPV33 L2 9 422 HPV45 E7 9 3										
HPV33 L2 9 422 HPV45 E7 9 3										
HPV33 L2 9 422										
HPV33 L2 8 338 HPV45 E7 10 3										
	HPV33	L2	8	338			nr v 4 3		10	,

Table XI HLA-B7 Supermotif Peptides

HPV45	E7	11	3	HPV45 L2	11	208
HPV45	L1	10	212	HPV45 L2	9	257
HPV45	L1	11	212	HPV45 L2	11	266
HPV45		8	386	HPV45 L2	10	94
HPV45		8	469	HPV45 L2	11	94
HPV45		9	469	HPV45 L2	8	445
HPV45		11	469	HPV45 L2	10	445
HPV45	L1	8	267	HPV45 L2	11	223
HPV45		9	267	HPV45 L2	8	97
HPV45		11	267	HPV45 L2	10	97
HPV45		8	21	HPV45 L2	11	97
HPV45		9	21	HPV45 L2 HPV45 L2	8	85
HPV45		10	21 .	HPV45 L2	9	85
HPV45		8	511	HPV45 L2	8	244
HPV45		8	338	HPV45 L2	9	452
HPV45		9	104	HPV45 L2	10	452
HPV45		9	39	HPV45 L2	11	452
HPV45		10	39	HPV45 L2	8	408
HPV45		8	119	HPV45 L2	9	377
HPV45		9	119	HPV45 L2	9	159
HPV45		10	119	HPV45 L2	10	159
HPV45		11	73	HPV45 L2	11	253
HPV45		8	464	HPV45 L2	10	355
HPV45		11	464	HPV45 L2	8	28
HPV45		9	213	HPV45 L2	11	354
HPV45		10	213	HPV45 L2	8	89
HPV45		8	437	HPV45 L2	11	284
HPV45		9	437	HPV45 L2	9	88
HPV45		9	40	HPV45 L2	9	324
HPV45		8	438	HPV45 L2	11	324
HPV45		11	438	HPV45 L2	10	419
HPV45		8	41	HPV45 L2	11	419
HPV45		10	439	HPV45 L2	11	137
HPV45		8	208	HPV45 L2	9	213
HPV45		10	208	HPV45 L2	- 9	360
HPV45		9	515	HPV45 L2	11	360
HPV45		8	525	HPV45 L2	9	184
HPV45		8	31	HPV45 L2	9	144
HPV45		9	31	HPV45 L2	9	271
HPV45		11	507	HPV45 L2	10	271
HPV45		8	321	HPV45 L2	9	398
HPV45		9	463	HPV45 L2	10	398
HPV45		8	189	HPV45 L2	11	398
HPV45		9	189	HPV45 L2	10	72
HPV45		9	89	HPV45 L2	11	72
HPV45		11	89	HPV45 L2	9	390
HPV45		9	247	HPV45 L2	11	390
HPV45		10	381	HPV45 L2	11	170
HPV45		9	436	HPV45 L2	10	119
HPV45		10	436	HPV45 L2	9	415
HPV45		10	79	HPV45 L2	8	370
HPV45		11	79	HPV56 E2	11	138
HPV45		9	258	HPV56 E2	10	243
HPV45		8	492	HPV56 E2	9	229
HPV45		9	492	HPV56 E2	10	229
HPV45		11	492	HPV56 E2	8	300
HPV45		9	27	HPV56 E2	9	300
HPV45		8	100	HPV56 E2	8	182
HPV45		8	208	HPV56 E2	9	186
13		-				

Table XI HLA-B7 Supermotif Peptides

				-				
HPV56	E2	9	151		HPV56		11	465
HPV56	E2	9	2		HPV56			40
HPV56	E2	11	2		HPV56	L1		40
HPV56		8 .	61		HPV56	L1		440
HPV56		10	61		HPV56	L1		196
HPV56		11	61		HPV56	L1	8	227
HPV56		8	24		HPV56	L1	9	227
HPV56		11	24		HPV56	L1	11	328
HPV56		9	7		HPV56	L1	8	51
HPV56		8	11		HPV56	L1	9	265
HPV56		11	11		HPV56	L1	11	265
HPV56		10	111		HPV56	L2	9	271
HPV56		8	46		HPV56	L2	10	271
		8	16	4	HPV56	L2	11	271
HPV56		8	63		HPV56		9	358
		9	63		HPV56		10	358
HPV56		10	63		HPV56		11	391
HPV56			5		HPV56		11	170
HPV56		8	5		HPV56		8	223
HPV56		9			HPV56		9	223
HPV56		11	5		HPV56		9	27
HPV56		9	521		HPV56		9	238
HPV56		11	219		HPV56		11	238
HPV56		8	472		HPV56		8	273
HPV56		9	472		HPV56		9	273
HPV56		11	472		HPV56		10	273
HPV56		8	11		HPV56		8	100
HPV56		10	11		HPV56		8	208
HPV56		11	11		HPV56		11	208
HPV56		8	318		HPV56		8	412
HPV56		9	318		HPV56		8	97
HPV56	L1	10	318				10	97
HPV56		8	30		HPV56		11	97
HPV56	L1	10	30		HPV56		8	215
HPV56		8	495		HPV56		9	215
HPV56	L1	9	495		HPV56			195
HPV56	L1	11	495		HPV56		8	119
HPV56	L1	9	96		HPV56		10	453
HPV56	L1	11	96		HPV56		10	453
HPV56	L1	8	343		HPV56		11	376
HPV56	L1	9	111		HPV56		8	373
HPV56	L1	9	48		HPV56		8	
HPV56	L1	10	48		HPV56		10	373 373
HPV56	L1	11	48		HPV56		11	
HPV56	L1	8	126		HPV56		8	409
HPV56	L1	9	126		HPV56		10	409
HPV56	L1	10	126		HPV56		11	409
HPV56	L1	10	220		HPV56		11	253
HPV56	L1	9	12		HPV56		9	159
HPV56	L1	10	12		HPV56		10	159
HPV56	L1	11	12		HPV56		8	89
HPV56	L1	8	319		HPV56		10	335
HPV56	L1	9	319		HPV56		11	284
HPV56		8	320		HPV56		8	244
HPV56		9	466		HPV56		9	88
HPV56		10	466		HPV56		8	77
HPV56		8	49		HPV56		11	77
HPV56		9	49		HPV56	L2	9	324
HPV56		10	49		HPV56		11	324
HPV56		11	441		HPV56		11	266
HPV56		10	465		HPV56	L2	9	422

Table XI HLA-B7 Supermotif Peptides

HPV56	L2	10	422
HPV56	L2	11	422
HPV56	L2	8	85
HPV56	L2	9	85
HPV56	L2	9	399
HPV56	L2	10	399
HPV56	L2	11	399
HPV56	L2	9	213
HPV56	L2	10	213
HPV56	L2	11	213
HPV56	L2	8	368
HPV56	L2	9	184
HPV56	L2	9	144
HPV56	L2	10	72
HPV56	L2	11	72
HPV56	L2	8	419
HPV56	L2	9	416
HPV56	L2	11	416

Table XI A. HPV 6A HLA-B7 Supermotif Peptides

			 ·				
2	3	4			E1	9	478
L2	9	271			E1	10	478
L2	10	271			L2	8	206
L2	8	118			E2	10	257
E1	10	337			E2	11	257
E1	11	337			L2	10	425
E2	8	326			L2	11	425
E2	10	326			L1	10	117
L2	10	410			L2	8	188
L1	8	158			L2	11	188
E4	11	39			L2	11	336
L2	8	217			L2	10	362
L2	9	217			L2	11	362
L2	8	98			L1	9	46
L2	10	98			E1	8	301
L2	11	98			E2	10	208
L2	9	182			E1	9	455
E1	10	181			E1	10	455
E5	9	77			E1	11	455
E5	10	77			L2	9	32
E5	11	77			L2	10	32
L2	9	27			L2	8	193
L1	10	181			L2	8	450
L1	11	181			L2	9	450
E1	10	560			L2	10	450
L2	11	236			E2	9	352
L2	8	101			E2	11	352
Ll	8	89			L1	8	432
L1	9	89		•	L1	10	432
L1	10	89			L1	11	432
E7	8	19			E6	9	109
E7	11	- 19			L2	9	79 79
L1	8	236			L2	10	305
L1	9	236			L1	8	74
L1	10	236			L1 L1	8 9	74
Ll	11	236				8	422
L1	8	434			L2	8	18
L1	9	434			E5 E5	9	18
L1	11	434				11	18
L2	11	354			E5 E1	11	591
E1	9	257			L1	8	134
E1	11	257			L1	10	134
E1	9	309			E1	11	544
E1	10	309			L1	8	390
E1	11	309			Li	10	390
L2	11	162			L2	10	358
L2	10	95 95			L2	9	157
L2	11				L1	8	15
L1	8	265			Li	11	15
L1	9 10	265 90			L2	9	251
E2	9	593			L2	11	251
E1	10	593			L2	9	123
E1	11	593 593			E7	10	17
E1		407			E1	8	479
L2 L2	10 9	402			E1	9	479
L2	11	402			E1	11	479
L2	8	85			L2	8	28
L2 L2	9	85			E4	8	28
	10	85			E1	8	310
ь2	10	0.5					

Table XI A. HPV 6A HLA-B7 Supermotif Peptide

E1	9	310
E1	10	310
E1	11	310
L1	9	182
L1	10	182 60
E2 E1	8 9	561
E1	11	561
L1	10	404
L1	8	13
L1	9	13
L1 E4	10 8	13 32
L1	11	403
L1	9	12
L1	10	12
L1	11	12 235
E2 E2	9	353
E2	10	353
E2	11	353
L2	9	170
L2	10	170
L2 L2	11 8	170 90
L2	9	90
E1	10	527
E1	11	527
E1	9	118
E1 E7	10 9	118 46
E7	10	46
E1	8	512
E1	9	512
E1	10	512
E1 E7	11 8	512 16
E7	11	16
L2	10	169
L2	11	169
L2	8	167
L2	8 10	381 381
L2 L2	10	284
L2	11	284
L1	9	481
L1	10	481
E1 E4	11 9	225 31
L2	9	89
L2	10	89
L2	10	219
L2	11	219
L1	8	3
L1 E2	9	246
E2	9	246
E2	11	246
L2	10	324
E1	9 -	93

D: J		
Peptid	ies	
E1	10	93
E2	9	323
E2	.10	323
E2	11	323
L1	8	402
E1	9	107
L2	11	417
E1	10	89
E1	11	89
L2	9	138
E2	9	215
E4	8	62
E4	11	62
Ll	10	427
L2	8	414
L2	9	414
E1	8	468
E1	9	468
E1	10	468
E5	8	63
E5	10	63
E5	11	63
L2	9	375
E4	9	27
E2	10	234
L2	8	242
L1	11	290
Ll	11	174
E2	11	221
L2	10	213
E1	9	641
L1	9	59
L1	11	59
L2	11	72 388
L2	10	388
L2	11	
L2	8	122 122
L2	10	
E2	9	59 11
L1	10	
L1	11 9	11 66
E4 L1	8	227
L1	9	227
L1	8	457
L1	9	457
E6	10	59
	1168107 vl	33
3.		

Table XI B. HPV 6B HLA-B7 Supermotif Peptides

				-			
2	3	4			L2	9	401
L2	9	271			L2	11	401
L2	10	271			L2	8	85
L2	8	118			L2	9	85
E1	10	337			L2	10	85
E1	11	337			E1	9	478
E2	. 8	326			E1	10	478
E2	10	326			L2	8	206
L2	10	409			E2	10	257
E4	8	3			E2	11	257
E4	9	3			L2	10	425
E4	10	3			L2	11	425
E4	11	3			L1	10	117
L1	8	158			L2	8	188
E4	11	49			L2	11	188
L2	8	217			L2	11	336
L2	9	217			L2	10	362
L2	8	98			L2	11	362
L2	10	98			Ll	9	46
L2	11	98			E1	8	301
E1 .	10	181			E2	10	208
L2	9	182			E1	9	455
E5A	8	77			E1	10	455
E5A	9	77			E1 L2	11 9	455 32
E5A	10	77			L2	10	32
E5A	11	77			L2	8	193
L2	9	27			L2	8	450
L1	10	181			L2	9	450
L1	11	181			L2	10	450
E1	10	560 236			E2	9	352
L2	11	101			E2	11	352
L2 L1	8	89			L1	8	432
L1	9	89			Ll	10	432
L1	10	89			L1	11	432
E7	8	19			E6	9	109
E7	11	19			L2	9	79
Li	8	236			L2	10	79
L1	9	236			Ll	8	305
L1	10	236			Ll	8	74
L1	11	236			L1	9	74
L1	8	434			L2	8	422
L1	9	434			E5A	8	18
L1	11	434			E5A	9	18
L2	11	354			E5A	11	18
El	9	257			E1	11	591
E1	11	257			L1	8	134
El	9	309			L1	10	134
El	10	309			E1	11	544 390
E1	11	309			L1 L1	8 10	390
L2	11	162			L2	10	358
L2	10	95			L2	9	157
L2	11	95			L1	8	15
L1	8	265			L1	11	15
L1	9	265			L2	9	251
E2	10	90			L2	11	251
E1	9	593			L2	9	123
E1	10	593			E7	10	17
E1 L2	11 10	593 406			E1	8	479
LL2	10	400					

Table XI B. HPV 6B HLA-B7 Supermotif Peptides

			HLA-B7 Supermotif F	eptides		
E1	9	479		E2	8	246
E1	11	479		E2	9	246
L2	8	28		E2	11	246
E4	8	38		L2	10	324
E1	8	310		E1	9	93
E1	9	310		E1	10	93
E1	10	310		E4	8	68
E1	11	310		E4	9	68
L1	9	182		L1	8	402
L1	10	182		E1	9	107
E2	8	60		E1	10	89 89
E1	9	561		E1 L2	9	138
E1	11	561		L2	9	419
L1	10	404 13		L2	10	419
L1	8 9	13		L2	11	419
L1 L1	10	13		E2	8	215
E4	8	42		E2	9	215
L1	11	403		E4	8	72
Li	9	12		E4 *	11	72
L1	10	12		L1	10	427
L1	11	12		L2	9	413
E2	9	235		L2	10	413
E2	8	353		E1	8	468
E2	10	353		E1	9	468
E2	11	353		E1	10 8	468 63
L2	9	170		E5A E5A	10	63
L2	10	170		E5A	11	63
L2	11	170 90		L2	9	375
L2 L2	8 9	90		E4	9	37
E1	10	527		E2	10	234
E1	11	527		L2	8	242
E1	9	118		L1	11	290
E1	10	118		L1	11	174
E7	9	46		L2	10	213
E7	10	46		E1	9	641
E1	8	512		L1	9	59
E1	9	512		L1	11	59 72
E1	10	512		L2	11 9	388
E1	11	512		L2 L2	10	388
E7	8	16		L2	11	388
E7	11	16 169		L2	8	122
L2 L2	10 11	169		L2	10	122
L2	8	167		E2	9	59
L2	8	381		L1	10	11
L2	10	381		L1	11	11
L2	10	284		E4	9	76
L2	11	284		L1	8	227
L1	9	481		L1	9	227
L1	10	481		L1	8	457
E1	11	225		L1	9 10	457 59
E4	9	41		E6	10	עכ
L2	9	89				
L2	10	89				
L2	10	219 219				
L2	11 8	3				
L1 L1	9	3				
	-	-				

Table XI C. HPV11 HLA-B7 Supermotif Peptides

2 L2 L2	3 	4	L2	10
	9			
т э		270	L2	9
	10	270	L2	11
E1	10	337	E1	10 .
E1	11	337	E2	8
E2	8	325	E2	10
E2	10	325	E2	11
L1	8	159	L2	8
	11	50	L2	11
E4		97	L2	10
L2	8	97	L2	11
L2	10		Ll	10
L2	11	97		
L2	9	181	L2	11
E4	9	47	L1	9
L2	9	26	. L2	9
L1	10	182	E1	8
Ll	11	182	E1	9
E1	10	560	E1	10
L2	11	235	E1	11
L2	11	353	L2	9
L2	8	100	L2	10
Li	8	89	ъ2	8
Li	9	89	L2	8
L1	10	89	L2	9
	8	19	L2	10
E7	11	19	L2	10
E7		357	L2	11
L2	11		E2	9
L1	8	237	E2	11
L1	9	237	L2	9
Ll	10	237		
Ll	11	237	L2	10
L1	8	435	E6	9
L1	9	. 435	L2	10
Ll	11	435	L1	8
L2	10	350	Ll	8
E1	9	257	L1	9
E1	11	257	L2	10
E1	9	309	L2	8
E1	11	309	ES	8
L2	11	161	E5	9
E2	8	210	E5	11
L2	10	94	E1	11
L2	11	94	L1	8
L1	8	266	Ll	10
Ll	9	266	L2	9
E2	10	90	L2	10
E5	8	77	E1	11
E5	9	77	L1	8
E5	10	77	Ll	10
		77	L1	8
E5	11		L1	11
E6	10	59		9
E1	9	593	L2	
E1	10	593	L2	11
E1	11	593	E4	8
Ll	8	458	E7	10
L1	9	458	E1	9
L2	8	216	El	11
	9	216	L2	8
			E4	8
L2 L2	10	402	P-4	
ь2 ь2 ь2	10 8	402 84	E1 E1	8 10

Table XI C. HPV11 HLA-B7 Supermotif Peptides

			rila-B/ Superii	iour Peptides	•
E1	11	310		L2	10
		183		E1	10
L1	9			E1	11
L1	10	183		L2	9
E1	9	561		L2	9
E1	11	561		L2 L2	10
E4	8	4.3		L2	
L1	10	405			11
Ll	8	13		E2 E4	10 11
L1	9	13			10
L1	10	13		L1	9
E4	9	42		L2 L2	10
L1	11	404		E1	8
L1	9	12		E1	9
L1	10	12		E1	10
Ll	11	12		E5	8
E2	8	352		E5	10
E2	10	352		L2	9
E2	11	352		E4	9
L2	9	122		L2	8
L2	9	169		L1	11
L2	10	169		E1	9
L2	11	169		E4	8
L2	8	89		E4	9
L2	9	89		E4	10
E1	10	527		E4	11
E1	11	527		L1	9
E1	9	118		L1	11
E1	10	118		L1	9
E1	8	512		L2	9
E1	9	512		L2	10
E1	10	512		L2	11
E1	11	512 46		L1	10
E7	9	46		Li	11
E7	10 8	16		L2	8
E7	11	16		L2	10
L2	10	168		ES	8
L2	11	168		E4	9
L2	8	166		L1	8
L2	10	283		L1	9
L2	11	283			
Li	10	482			
E1	11	225			
E4	8	67			
E4	9	67			
E4	10	41			
L2	9	88			
L2	10	88			
L2	10	218			
L2	11	218			
L1	8	3			
L1	9	3			
E2	9	195			
E2	9	322			
E2	10	322			
E2	11	322			
E1	9	93			
B1	10	93			
L2	11	76			
Li	8	403			
	-				

E1 9

107-

Table XII HLA-B27 Supermotif Peptides

						_	
1	2	3	4	HPV16		8	446
HPV16		10	67	HPV16		9	446
HPV16	E1	9	203		E1	11	446
HPV16	E1	9	209	HPV16		8	491
HPV16	E1	11	209	HPV16		9	491
HPV16	E1	9	516	HPV16	E1	11	491
HPV16	E1	11	516	HPV16	E1	8	229
HPV16	E1.	8	399	HPV16	E1	10	229
HPV16		10	399	HPV16	E1	11	229
HPV16		8	76	HPV16	E1	8	286
HPV16		10	76	HPV16	E1	9	286
HPV16		11	76	HPV16		8	581
HPV16		9	627 .	HPV16	E1	8	468
HPV16		8	606	HPV16		9	468
HPV16		9	431	HPV16		11	468
HPV16		10	416	HPV16		8	310
			416	HPV16		9	310
HPV16		11		HPV16		10	310
HPV16		8	26	HPV16		11	310
HPV16		8	291	HPV16		8	108
HPV16		10	291			8	78
HPV16		11	116	HPV16			78
HPV16		10	182	HPV16		9	347
HPV16		9	211	HPV16		9	
HPV16		11	226	HPV16		11	347
HPV16	E1	10	284	HPV16		10	341
HPV16	E1	11	284	HPV16		11	150
HPV16	E1	9	482	HPV16		9	183
HPV16	E1	10	482	HPV16		11	410
HPV16	E1	8	79	HPV16		В	88
HPV16	E1	8	251	HPV16		11	88
HPV16	E1	10	251	HPV16		11	124
HPV16	E1	11	251	HPV16		8	506
HPV16	E1	8	426	HPV16	E1	9	506
HPV16	E1	8	550	HPV16	E1	11	506
HPV16	E1	8	469	HPV16	E1	9	295
HPV16	E1	10	469	HPV16	E1	11	295
HPV16	E1	11	469	HPV16	E1	10	504
HPV16		9	417	HPV16	E1	11	504
HPV16		10	417	HPV16	E1	8	119
HPV16		8	413	HPV16	E1	9	119
HPV16		10	413	HPV16	E1	9	614
HPV16		9	87	HPV16	E1	10	574
HPV16		10	262	HPV16	E1	11	574
HPV16		8	579	HPV16	E1	11	402
HPV16		10	579	HPV16		9	549
HPV16		8	557	HPV16		8	439
HPV16		8	460	HPV16		10	439
HPV16		10	86	HPV16	E1	9	609
HPV16		9	393	HPV16	E1	10	281
HPV16		10	393	HPV16	E1	9	412
		8	198	HPV16		11	412
HPV16				HPV16		8	322
HPV16		9	198	HPV16		11	322
HPV16		10	198	HPV16		9	46
HPV16		11	198	HPV16		9	28
HPV16		9	536			11	28
HPV16		10	536	HPV16			90
HPV16		8	312	HPV16		8	90
HPV16		9	312	HPV16		10	
HPV16		10	312	HPV16		8	176
HPV16	E1	11	312	HPV16	E2	9	176

Table XII HLA-B27 Supermotif Peptides

			•	-		
HPV16	E2	11	176	HPV16 E2	8	213
HPV16		8	171	HPV16 E2	9	213
HPV16		8	51	HPV16 E2	9	342
HPV16		9	51	HPV16 E2	11	342
HPV16		8	305	HPV16 E2	10	6
		10	305	HPV16 E2	11	6
HPV16				HPV16 E2	8	65
HPV16		11	305	HPV16 E2	9	65
HPV16		9	323			65
HPV16		10	323	HPV16 E2	11	
HPV16	E2	11	323	HPV16 E2	8	179
HPV16	E2	8	328	HPV16 E2	10	179
HPV16	E2	10	328	HPV16 E2	9	135
HPV16	E2	8 -	258	HPV16 E2	11	135
HPV16	E2	10	258	HPV16 E2	8	222
HPV16	E2	11	258	HPV16 E2	10	17
HPV16	E2	8 .	110	HPV16 E2	9	254
HPV16		10	110	HPV16 E2	8	186
HPV16		10	211	HPV16 E2	9	186
HPV16		11	211	HPV16 E2	8	160
HPV16		8	164	HPV16 E2	9	160
		8	307	HPV16 E2	10	160
HPV16		9	307	HPV16 E2	10	289
HPV16		-		HPV16 E2	8	326
HPV16		8	112		10	326
HPV16		10	112	HPV16 E2		
HPV16	E2	8	52	HPV16 E2	9	350
HPV16	E2	11	52	HPV16 E2	8	319
HPV16	E2	9	327	HPV16 E2	9	33
HPV16	E2	11	327	HPV16 E2	10	33
HPV16	E2	8	34	HPV16 E2	11	33
HPV16	E2	9	34	HPV16 E2	8	44
HPV16	E2	10	34	HPV16 E2	11	44
HPV16	E2	11	34	HPV16 E2	9	303
HPV16		9	306	HPV16 E2	10	303
HPV16		10	306	HPV16 E5	8	57
HPV16		9	111	HPV16 E5	9	57
HPV16		11	111	HPV16 E5	10	57
HPV16		9	257	HPV16 E5	11	57
HPV16		11	257	HPV16 E5	8	74
		8	298	HPV16 E5	9	74
HPV16			291	HPV16 E5	8	29
HPV16		8		HPV16 E5	11	29
HPV16		11	291	HPV16 E6	10	40
HPV16		8	26		11	40
HPV16		11	26		10	71
HPV16		11	301	HPV16 E6		
HPV16		9	129	HPV16 E6	9	127
HPV16		10	129	HPV16 E6	9	14
HPV16		11	129	HPV16 E6	8	132
HPV16	E2	8	36	HPV16 E6	8	54
HPV16	E2	9	36	HPV16 E6	8	137
HPV16	E2	11	36	HPV16 E6	9	30
HPV16	E2	10	217	HPV16 E6	10	30
HPV16	E2	11	217	HPV16 E6	10	108
HPV16		8	55	HPV16 E6	10	135
HPV16		9	55	HPV16 E6	8	128
HPV16		10	55	HPV16 E6	10	74
HPV16		9	67	HPV16 E6	8	45
HPV16		11	67	HPV16 E6	9	45
HPV16		8	181	HPV16 E6	10	45
				HPV16 E6	8	1
HPV16		10	353	HPV16 E6	9	î
HPV16	E2	11	353		-	-

Table XII HLA-B27 Supermotif Peptides

				•	•			
HPV16	E6	8	100		HPV16	L1	8	79
HPV16		9	100		HPV16			79
HPV16	E6	11	16		HPV16	L1		79
HPV16	E6	10	130		HPV16			468
HPV16		10	123		HPV16			468
HPV16		9	124		HPV16	L1	9	500
HPV16		8	84		HPV16	L1	9	477
HPV16		10	17		HPV16	Ll	9	467
HPV16		8	46		HPV16	L1	11	467
HPV16		9	46		HPV16	L1	9	390
HPV16		9	78		HPV16	L1	11	390
HPV16		11	78		HPV16	L1	8	276
HPV16		9	150		HPV16	L1	11	276
HPV16		8	61		HPV16	L1	10	107
HPV16		9	61		HPV16	L1	11	107
HPV16		8	83		HPV16	L1	8	84
HPV16		9	83		HPV16	L1	11	84
HPV16		8	50		HPV16	L1	9	150
HPV16		9	59		HPV16	L1	11	150
HPV16		11	59		HPV16	L1	9	334
HPV16		8	4.8		HPV16	L1	9	242
HPV16		10	4.8		HPV16	L1	10	288
HPV16		8	76		HPV16	L1	9	169
HPV16		9	76		HPV16	L1	9	462
HPV16		8	8		HPV16	L1	11	462
HPV16		10	65		HPV16	L1	10	504
HPV16		8	1		HPV16	L1	8	89
HPV16		11	1		HPV16	L1	10	89
HPV16		8	72		HPV16	L1	11 '	89
HPV16		11	72		HPV16	L1	10	340
HPV16		8	524		HPV16	L1	8	122
HPV16		9	55		HPV16	L1	10	122
HPV16		9	405		HPV16	L1	8	391
HPV16		10	405		HPV16	L1	10	391
HPV16		9	187		HPV16		11	391
HPV16		9	255		HPV16	L1	9	284
HPV16		10	255		HPV16	L1	10	284
HPV16		10	479		HPV16		9	492
HPV16	L1	11	386		HPV16		11	492
HPV16	L1	11	99		HPV16		10	277
HPV16	L1	8	344		HPV16		11	277
HPV16	L1	8	145		HPV16		9	45
HPV16	L1	9	196		HPV16		10	45
HPV16	L1	10	196		HPV16		11	45
HPV16	L1	11	196		HPV16		10	66
HPV16	L1	8	134		HPV16		11	66
HPV16	L1	10	134		HPV16		8	363
HPV16		8	491		HPV16		10	363
HPV16	L1	10	491		HPV16		11	363
HPV16	Ll	9	101		HPV16		8	283
HPV16		11	101		HPV16		10	283
HPV16		8	417		HPV16		11	283
HPV16		9	417		HPV16		8	61
HPV16	L1	8	303		HPV16		9	61
HPV16		10	303		HPV16		10	61
HPV16		11	303		HPV16		11	61
HPV16	L1	9	78		HPV16		9	21
HPV16	L1	10	78		HPV16		10	21
HPV16		11	78		HPV16		11	21
HPV16	Ll	8	261		HPV16	ьī	9	381

Table XII HLA-B27 Supermotif Peptides

				TIEA-BET Supermont replace	•		
HPV16 I	.1 :	10	381	HPV1	5 L2	11	314
HPV16 I	.1	В	177	HPV1	5 L2	10	324
HPV16 I	1 :	9	177	HPV1		8	235
HPV16 I	1 (8	444	HPV1	5 L2	9	235
HPV16 I	.1 :	9	444	HPV1		10	89
HPV16 I	1 :	10	444	HPV1		10	269
HPV16 I	1 1	8	96	HPV1		8	578
HPV16 I	.2		7	HPV1:		9	578
HPV16 I	.2	9	322	HPV1:		8	298
HPV16 I			22	HPV1:		10	298
HPV16 I		10	179	HPV1:		8	125
HPV16 I			317	HPV1		8	406
HPV16 I		10	317	HPV1:		10	406
		11	317	HPV1		8	276
HPV16 I		8	38	HPV1		9	276
		9	38	HPV1		10 11	276 276
HPV16 I		11	38	HPV1:		8	25
		8	68	HPV1:		8	602
		11	3	HPV1		10	602
HPV16 I		8	11	HPV1		8	613
		9	11	HPV1		8	236
HPV16 I		10	460	HPV1		9	236
HPV16 I		8	457 457	HPV1		11	236
HPV16 I		9	457	HPV1		8	555
HPV16 I		10 10	45/	HPV1		10	555
HPV16 I		8	8	HPV1		9	101
		11	8	HPV1		9	621
HPV16 I		8	448	'HPV1		10	70
HPV16 I		8	455	HPV1		9	218
HPV16		10	455	HPV1		11	233
		11	455	HPV1	8 E1	8	258
		9	312	HPV1	8 E1	9	258
HPV16		11	312	HPV1	8 E1	10	258
HPV16	L2	8	34	HPV1	8 E1	11	258
		9	34	HPV1		8	291
HPV16	L2	8	459	HPV1	B E1	11	291
HPV16		11	459	HPV1		9	489
HPV16		9	456	HPV1		10	489
HPV16	L2	10	456	HPV1		8	498
HPV16	L2	11	456	HPV1		9	498
HPV16		8	458	HPV1		11	498
HPV16		9	458	HPV1		9	575
HPV16		10	297	HPV1		11	575
		9	353	HPV1		8	433
		9	304	HPV1		8	557 417
		9	219	HPV1		11 8	123
		10	219	HPV1 HPV1		10	123
HPV16		8	296	HPV1		11	476
		11	296	HPV1 HPV1		9	424
		9	229	HPV1		10	424
		10	229	HPV1		8	629
HPV16		9	190	HPV1		8	82
		10	247	HPV1		9	82
		11	247	HPV1		8	305
		9	10	HPV1		9	305
		10	10	HPV1		10	305
		10	77 77	HPV1		8	564
	L2	9	314	HPV1		11	409
HPV16	112	,	214	11171			

Table XII HLA-B27 Supermotif Peptides

				THE TOTAL	Juponnoni i e	Pudeo			
HPV18	E1	10	471		H	IPV18	E2	11	73
HPV18	E1	11	471			IPV18		9	237
HPV18	E1	9	400		H	IPV18	E2	10	237
HPV18	E1	10	400			IPV18		9	50
HPV18	E1	9	205			IPV18		10	21
HPV18	E1	10	205				E2	10	311
HPV18	E1	11	205			PV18		8	295
HPV18	E1	10	648			IPV18		10	295
HPV18	E1	8	319			IPV18		11	52
HPV18	E1	9	319			(PV18		8	256
HPV18		10	319			IPV18		11	256
HPV18		11	319			IPV18		8	328
HPV18	E1	10	286			IPV18		10	328
HPV18		8	453			IPV18		9	181
HPV18		9	453			IPV18		11	181
HPV18		11	453			IPV18		10	10
HPV18		10	543			IPV18		11	
HPV18		8	209			IPV18		8	114
HPV18		9	209			IPV18		10	114
HPV18		10	209			IPV18		11	176
HPV18		10	581			IPV18		8	74
HPV18		8	475			IPV18		9	74
HPV18		8	317			IPV18		10	74
HPV18		9	317			IPV18		11	159
HPV18		10	317			IPV18		10	290
HPV18		11	317			IPV18		8	40
HPV18 HPV18		9	111 354			IPV18		9	40
HPV18		11	354			IPV18		10	40
HPV18		8	122			IPV18		11	40
HPV18		9	122			IPV18		9	308
HPV18		11	122			IPV18		9	115
HPV18		10	423			IPV18		10	115
HPV18		11	423		H	IPV18	E2	11	115
HPV18		10	348		H	IPV18	E2	8	299
HPV18		9	556		F	IPV18	E2	8	292
HPV18		8	420		H	IPV18	E2	11	292
HPV18	E1	10	420		F	IPV18	E2	8	306
HPV18	E1	11	127			PV18		11	306
HPV18	E1	8	513			IPV18		8	59
HPV18	E1	9	438			IPV18		10	59
HPV18	E1	8	588			IPV18		8	128
HPV18	E1	11	588			IPV18		9	128
HPV18	E1	9	293			IPV18		10	128
HPV18		9	523			IPV18		11	128
HPV18		8	75			IPV18		9	4
HPV18		10	75			IPV18		9 10	258 258
HPV18		11	75					11	222
HPV18		9	616			IPV18		9	342
HPV18		8	446			IPV18		10	342
HPV18		10	446			IPV18		11	342
HPV18		8	288			IPV18		10	307
HPV18		10	288			IPV18		11	279
HPV18		11	288			IPV18		8	158
HPV18		9	419 419			IPV18		9	71
HPV18		11	329			IPV18	E2	11	71
HPV18 HPV18		8 11	329			IPV18	E2	8	27
HPV18		TT	73			IPV18		10	27
HPV18		10	73			IPV18		11	27
HEVIO		-0			•	-			